The state of the s	** ***		00 1514077			10010000
Kelch domain confaming	VLUDC1					1557737 s at
solate carrier ranny 2 (racintated glucose trans	AL HIDOI	122772	NID 7510/12			1552733 at
solute carrier family 2 (facilitated changes tones	SLC2A13	114134	NP 443117	NM 052885	4D	1552695 a at
hypothetical protein FLJ22313	FLJ22313	64224	NP_071768	NM_022373	4D	1552628_a_at
tRNA nucleotidyl transferase, CCAadding, 1	TRNTI	51095	NP_057084	NM_016000	4D	1552625_a_at
ring finger and WD repeat domain 2	RFWD2	64326				1552617 a at
chromosome 6 open reading frame 141	C6orf141	135398	NP_699175			15525/5_a_at
caspase recruitment domain family, member 12	CARD12	58484	NP_067032	021209		
membe	SLC34A3	172000	111 070100			
raser syndrome 1	FRASI	147600				
Zinc ringer protein 31 (KOX 29)	EDASI	90144	116262			1552503 at
The second of th	ZNE21	7570	ND 660381	NM 145238	4D	1552499 a at
lactation elevated 1	LACE1	246269	NP 660358	NM_145315	4D 1	1552490_at
centaurin, beta 2	CENTB2	23527	NP_036419	NM_012287	4D	1552472_a_at
hypothetical protein LOC126248	LOC126248	126248	NP_775750	NM_173479	4D	1552452_at
kelchlike 10 (Drosophila)	KLHL10	317719	NP_689680	NM_152467	4D	1552425_a_at
hypothetical protein FLJ36119	FLJ36119	254173	NP_694986			1552419_s_at
phosphodiesterase 7A	PDE7A	5150				1552343_s_at
goosecoid	GSC	145258	NP_776248			155255 at
GTPase, IMAP family member 1	GIMAPI	170575		130759		13
microfibrillarassociated protein 3	MFAP3	4238	NP_005918	005927		l a
hypothetical protein MGC29937	MGC29937	123207	NP_653198	14439/		122210 at
phosphatidylinositol glycan, class X	PIGX	54965				1552210 ±
AFG3 ATPase family gene 3like 1 (yeast)	AFG3L1	172	NP_001123			155228/_s_at
ATPase, H+ transporting, lysosomal 31kDa, V1 subun	ATP6V1E2	90423	NP_542384			1552286_at
similar to RIKEN cDNA 5730528L13 gene	MGC17337	91283	NP_542386			15522//_a_at
PX domain containing serine/threonine kinase	PXK	54899	NP_060241			1552274_at
mitogenactivated protein kinase 1	MAPKI	5594	NP_620407			1552264_a_at
mitogenactivated protein kinase 1	MAPK1	5594	NP_620407			1552263_at
WAP fourdisulfide core domain 2	WFDC2	10406	NP_542773			1552261_at
hypothetical protein MGC4677	MGC4677	112597	NP_443103			
protein tyrosine phosphatase, nonreceptor type 21	PTPN21	11099	NP_008970			
paired box gene 8	PAX8	7849	NP_003457			121_at
	Gene Symbol	LocusLink	HumanProtAcces	HumanRNAAccession	ExperimentID	Clone ID

subun 126 hypothetical protein MGC14126 phosphodiesterase 5A, cGMPspecific RAS and EF hand domain containing zinc finger protein 441	ZNF441	126068	NP_689568	DIM COCCU	3 8	
				~ ア・ファ ーナ・フルナナ		11000195 at
	RASEF	158158	NP_689786	NM_152573	4D	
	RASEF	158158	NP_689786	NM_152573	40	1555185_at
	PDE5A	8654	NP_237223	NM_033431	40	
	MGC14126	84984		NM_032898	4D	a
	ATP6V0D2	245972	NP_689778	NM_152565	4D	
	FLJ25422	202151	NP_659437	NM_145000	4D	1553147_at
chromosome 9 open reading frame 72	C9orf72	203228	NP_659442	NM_145005	4D	11
RBassociated KRAB repressor	RBAK	57786	NP_066986	NM_021163	4D	1553122_s_at
nuclear transcription factor, Xbox binding 1	NFXI	4799	NP_667345	NM_147134	4D	1553103_at
tigger transposable element derived 1	TIGDI	200765	NP_663748	NM_145702	4D	1553099_at
	BCL2L11	10018	NP_619533	NM_138627	4D	1553096_s_at
BCL2like 11 (apoptosis facilitator)	BCL2L11	10018	NP_619532	NM_138626	4D	1553088_a_at
S. C.				NM_032362	4D	1553087_at
ankyrin repeat and SOCS boxcontaining 10	ASB10	136371	NP_543147	NM_080871	4D	1553039_a_at
synaptotagminlike 5	SYTL5	94122	NP_620135	NM_138780	4D	1553033_at
sulfite oxidase	SUOX	6821	NP_000447	NM_000456	4D	1553030_a_at
0 0				NM_054029	4D	1552955_at
mel transforming oncogenelike	MELL1	79258	NP_258428	NM_033467	4D	1552930_at
fidgetinlike I	FIGNLI	63979	NP_071399	NM_022116	4D	1552921_a_at
interleukin 23 receptor	IL23R	149233	NP_653302	NM_144701	4D	1552912_a_at
	C10orf27	219793	NP_689923	NM_152710	4D	1552887_at
	АТОН7	220202	NP_660161	NM_145178	4D	1552879_a_at
5hydroxytryptamine (serotonin) recentor 6	HTR6	3362	NP_000862	NM_000871	4D	1552857_a_at
	ZNF619	285267	NP_775927	NM_173656	4D	1552836_at
tolllike receptor 4	TLR4	7099	NP_612567	NM_138557	4D	1552798_a_at
				NM_153039	4D	1552789_at
helicase (DNA) B	HELB	92797	NP_387467	NM_033647	4D	1552788_a_at
	CLEC4D	338339	NP_525126	NM_080387	4D	1552772_at
, <u></u>	CAMKK	84254	NP_757344	NM_172207	4D	1552768_at
	MCF2L	23263	NP_079255	NM_024979	4D	1552743 at
ligase 2						

hypothetical protein FLJ25371	FLJ25371	152940	NP_689756	NM_152543	4D	1553838_at
SRY (sex determining region Y)box 3	SOX3	6658	NP_005625	NM_005634	4D	1553802_a_at
				NM_032202	4D	1553793_a_at
SLAM family member 9	SLAMF9	89886	NP_254273	NM_033438	4D	1553770_a_at
discoidin, CUB and LCCL domain containing 1	DCBLD1	285761	NP_775945	NM_173674	4D	1553768 a at
G proteincoupled receptor 97	GPR97	222487	NP_740746	NM_170776	4D	1553723_at
hypothetical protein FLJ90396	FLJ90396	163049	NP_699189	NM_153358	4D	1553704_x_at
hypothetical protein FLJ90396	FLJ90396	163049	NP_699189	NM_153358	4D	1553703_at
Sp1 transcription factor	SPI	6667	NP_612482	NM_138473	4D	1553685_s_at
putative MAPK activating protein	MGC3794	261726	NP_690866	NM_152902	4D	1553677_a_at
hypothetical protein FLJ40342	FLJ40342	124989	NP_689560	NM_152347	4D	1553626_a_at
sperm associated antigen	SPAG11	10407	NP_478113	NM_058206	4D	1553579 a at
interferon epsilon 1	IFNE!	338376	NP_795372	NM_176891	4D	1553574_at
histone 1, H1t	HISTIHIT	3010	NP_005314	NM_005323	4D	1553568 a at
vomeronasal 1 receptor 5	VNIR5	317705	NP_776257	NM_173858	4D	1553550_at
keratin 6 irs4	K6IRS4	121391	NP_778223	NM_175053	4D	1553539_at
NACHT, leucine rich repeat and PYD containing 8	NALP8	126205	NP_789781	NM_176811	4D	1003026_at
homeo box C12	HOXC12	3228	NP_776272	NM_173860	Đ	1553512_at
paralemmin 2	PALM2	114299	NP_443749	NM_053016	4D	1553502_a_at
hypothetical protein FLJ90834	FLJ90834	283135	NP_775849	NM_173578	4D	1553438_at
hypothetical protein FLJ31952	FLJ31952	146857	NP_653283	NM_144682	4D	1553423 a at
ataxin 2binding protein 1	A2BP1	54715	NP_665899	NM_145892	4D	1553422_s_at
hypothetical protein FLJ25168	FLJ25168	147081	NP_689679	NM_152466	4D	1553400_a_at
hypothetical protein FLJ32978	FLJ32978	128025	NP_653226	NM_144625	4D	1553373_at
chromosome 6 open reading frame 195	C6orf195	154386	NP_689767	NM_152554	4D	1553363_at
solute carrier family 18 (vesicular monoamine), me	SLC18A2	6571	NP_003045	NM_003054	#5	1000020_a_at
hypothetical protein FLJ33860	FLJ33860	284756	NP_775915	NM_173644	14	1553378 a at
colony stimulating factor 3 receptor (granulocyte)	CSF3R	1441	NP_758519	NM_172313	145	155324/ a_at
G proteincoupled receptor 128	GPR128	84873	NP_116176	NM_032787	4D	1553296_at
solute carrier family 36 (proton/amino acid sympor	SLC36A1	206358	NP_510968	NM_0/8483	40	12/2/2 al
hypothetical protein FLJ90036	FLJ90036	255403	NP_710158	NM_153691	40	1553269_at
Fanconi anemia, complementation group B	FANCB	2187	NP_689846	NM_152633	4D	1553244_at

transmembrane protein 37	TMEM37	140738	NP_899063	NM_183240	4D	1554483_at
PHD finger protein 20like 1	PHF20L1	51105	NP_079154	NM_024878	40	1554472_a_at
WINS1 protein with Drosophila Lines (Lin) homologo	WINSI	55180	NP_859524	NM_181740	£	
WINS1 protein with Drosophila Lines (Lin) homologo	WINSI	55180	NP_859523	NM_181739	40	1554455_at
neuroligin 2	NLGN2	57555	NP_065846	NM_020795	£ £	1554428 s at
anterior pharynx defective 1 homolog A (C. elegans	APHIA	51107	NP_057106	NM_016022	1	S
hypothetical protein FLJ20125	FLJ20125	54826	NP_060146	NM_017676	4D	1554345_a_at
DNA helicase HEL308	HEL308	113510	NP_598375	NM_133636	4D	1554342_s_at
histone deacetylase 4	HDAC4	9759	NP_006028	NM_006037	4D	1554322_a_at
hypothetical protein LOC136306	LOC136306	136306	NP_777619	NM_174959	4D	1554300_a_at
adult retina protein	LOC153222	153222	NP_705835	NM_153607	4D	1554229_at
PHD finger protein 21A	PHF21A	51317	NP_057705	NM_016621	4D	1554153 a at
unc51like kinase 2 (C. elegans)	ULK2	9706	NP_055498	NM_014683	4D	1554112 a_at
amyotrophic lateral sclerosis 2 (juvenile) chromos	ALS2CR16	130029	NP_991106	NM_205543	4D	1554108 at
DEAD (AspGluAlaAsp) box polypeptide 51	DDX51	317781	NP_778236	NM_175066	4D	1554085_at
serine palmitoyltransferase, long chain base subun	SPTLC1	10558	NP_847894	NM_178324	4D	1554053_at
zinc finger protein 12 (KOX 3)	ZNF12	7559	NP_008887	NM_006956	4D	1554021_a_at
apoptosisrelated protein PNASI	FLJ39616	51275	NP_057618	NM_016534	4D	1553987_at
chromosome 20 open reading frame [6]	C20orf161	90203	NP_690857	NM_152897	4D	1553960_at
UDPGal:betaGal beta 1,3galactosyltransferase polyp	B3GALT6	126792	NP_542172	NM_080605	4D	1553959_a_at
exosome component 6	EXOSC6	118460	NP_478126	NM_058219	4D	1553947_at
				NM_131915	4D	1553900_s_at
hypothetical protein FLJ40365	FLJ40365	126402	NP_775753	NM_173482	4D	1553893_at
G proteincoupled receptor MRGX2	MRGX2	117194	NP_473371	NM_054030	4D	1553889_at
hypothetical protein MGC35338	MGC35338	130619	NP_689602	NM_152389	4D	1553886_at
hypothetical protein MGC34713	MGC34713	285600	NP_775936	NM_173665	4D	1553868_a_at
G. C.				NM_138339	4D	1553865_a_at
chromosome 10 open reading frame 64	C10orf64	159491	NP_775795	NM_173524	4D	1553863_at
hypothetical protein MGC40368	MGC40368	255394	NP_689985	NM_152772	4D	1553861_at
Cohen syndrome	СОН1	157680	NP_689777	NM_152564	4D	1553852_at
chromosome 21 open reading frame 88	C21orf88	114041	NP_715635	NM_153754	4D	1553850_at

ST3 betagalactoside alpha2,3sialyltransferase 3	ST3GAL3	6487	NP_///624	NM_1/4964	40	1333/02_a_at
SLC2A4 regulator	SLC2A4KG	30/31	NF_004440	120007		د ا
Interieukin 28 receptor, alpha (interferon, lambda	ST COA AND	107702	NID 064446	NM 020062	4)	ا ا
	11 78B A	163702	NP 775088	NM 173065	4D	1555499 a at
cytochrome P450, family 4, subfamily B, polypeptid	CYP4B1	1580	NP_000770	NM_0007/9	4D	155549/_a_at
serologically defined colon cancer antigen 10	SDCCAG10	10283	NP_005860	NM_005869	3 5	1555495 a at
hypothetical gene supported by BC046200	LOC388407	388407	NP_982249	NM_203425	4D	at
transmembrane protein 1	TMEMI	7109	NP_001001723	NM_001001723	40	
Nacylsphingosine amidohydrolase (acid ceramidase)	ASAHI	427	NP_004306	NM_004315	40	a
FGD1 family, member 3	FGD3	89846	NP_149077	NM_033086	40	1555407_s_at
ATPbinding cassette, subfamily B (MDR/TAP), member	ABCB5	340273	NP_848654	NM_1/8559	40	10005/1_at
KIAA0377 gene product	KIAA0377	9677	NP_055474	NM_014659	4D	1555255_a_at
retinitis pigmentosa 1like 1	RPILI	94137	NP_849188	NM_178857	4D	1555245_s_at
hypothetical gene supported by BC055092	LOC401466	401466	XP_376783	XM_376783	4D	1555243_x_at
hypothetical gene supported by BC055092	LOC401466	401466	XP_376783	XM_376783	4D	1555241_at
four and a half LIM domains 5	FHL5	9457	NP_065228	NM_020482	4D	1555191_a_at
preBcell colony enhancing factor 1	PBEFI	10135	NP_877591	NM_182790	40	155516/_s_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_006766	NM_006775	4D	1555154_a_at
hypothetical protein MGC40069	MGC40069	348035	NP_872421	NM_182615	4D	1555073_at
scavenger receptor class F, member 1	SCARF1	8578	NP_663324	NM_145349	4D	1555021_a_at
zinc finger protein 398	ZNF398	57541	NP_733787	NM_170686	4D	1555015_a_at
hypothetical protein LOC403312	MGC39545	403312	NP_982277	NM_203452	4D	1555002_at
phospholipase A2, group IVD (cytosolic)	PLA2G4D	283748	NP_828848	NM_178034	4D	1554914_at
RAB39, member RAS oncogene family	RAB39	54734	NP_059986	NM_017516	4D	1554800_at
hypothetical protein FLJ32130	FLJ32130	146540	NP_689671	NM_152458	4D	1554770_x_at
hypothetical protein FLJ32130	FLJ32130	146540	NP_689671	NM_152458	4D	1554769_at
transforming, acidic coiledcoil containing protein	TACC1	6867	NP_006274	NM_006283	4D	1554690_a_at
hypothetical protein LOC126295	LOC126295	126295	NP_775751	NM_173480	4D	1554628_at
polypyrimidine tract binding protein 2	PTBP2	58155	NP_067013	NM_021190	4D	1554614_a_at
cyclin M2	CNNM2	54805	NP_060119	NM_017649	4D	1554523_a_at
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	4D	1554510_s_at
regulator of Gprotein signalling 7	RGS7	6000	NP_002915	NM_002924	4D	1554500_a_at

zinc finger protein 548	ZNF548	147694	NP_690873	NM_152909	40	1558622_a_at
zinc finger protein 621	ZNF621	285268	NP_940886	NM_198484	4D	1558620_at
LATI3TM protein	LAT13TM	81893	NP_112488	NM_031211	4D	1558534_at
hypothetical protein LOC113230	LOC113230	113230	XP_053966	XM_053966	4D	1558412_at
hypothetical gene supported by NM 194304	LOC439911	439911	XP_498467	XM_498467	4D	1558545_a_at
chromosome 22 open reading frame 15	C22orf15	150248	NP_872326	NM_182520	4D	1558333_at
nuclear factor of kappa light polypeptide gene enh	NFKBIL2	4796	NP_038460	NM_013432	4D	1558330_x_at
zinc finger, DHHC domain containing 18	ZDHHC18	84243	NP_115659	NM_032283	4D	1 22
phosphatidylinositol glycan, class W	PIGW	284098	NP_848612	NM_178517	4D	1558292_s_at
SFRS protein kinase 2	SRPK2	6733	NP_872633	NM_182691	4D	S
male sterility domain containing 2	MLSTD2	84188	NP_115604	NM_032228	4D	1558014_s_at
hypothetical protein FLJ21908	FLJ21908	79657	NP_078880	NM_024604	4D	1557984 s_at
catenin (cadherinassociated protein), delta 1	CTNNDI	1500	NP_001322	NM_001331	4D	1557944 s_at
chr3 synaptotagmin	CHR3SYT	83850	NP_114119	NM_031913	4D	1557793_at
phosphatidylinositol3phosphate/phosphatidylinosit o	PIP5K3	2005/6	NP_001002881	NM_001002881	+	100//17_dt
hypothetical gene supported by AK093266	LOC400238	400238	XP_378473	XM_3/84/3	3 8	155/05/_a_at
zinc finger protein 230	ZNF230	7773	NP_006291	NM_006300	40	at
NIMA (never in mitosis gene a) related kinase 8	NEK8	284086	NP_835464	NM_178170	4D	155/1/2_x_at
kelchlike 18 (Drosophila)	KLHL18	23276	NP_079286		4D	1557165_s_at
hypothetical protein FLJ22374	FLJ22374	84182	NP_115598	NM_032222	4D	1557141_at
hypothetical protein FLJ36070	FLJ36070	284358	NP_872380	NM_182574	4D	1557091_at
hypothetical protein LOC133491	LOC133491	133491	XP_376444	XM_376444	4D	1557056_at
similar to Zinc finger protein Rlf (Rearranged Lmy	LOC440971	440971	XP_496651	XM_496651	40	1000/44_a_at
hypothetical protein LOC283687	LOC283687	283687	NP_787094	NM_175898	4D	1556588_at
hypothetical protein LOC197322	LOC197322	197322	NP_777577	NM_174917	4D	1556552_a_at
FGFR1 oncogene partner 2	FGFR1OP2	26127	NP_056448	NM_015633	4D	1556283 s_at
chromosome 10 open reading frame 90	C10orf90	118611	NP_001004298	NM_001004298	4D	1556200_a_at
peroxisome biogenesis factor 13	PEX13	5194	NP_002609	NM_002618	4D	1556009_at
histidine triad nucleotide binding protein 1	HINTI	3094	NP_005331	NM_005340	4D	1555961_a_at
cell division cycle 25A	CDC25A	993	NP_001780	NM_001789	4D	1555772_a_at
angiotensin II receptorassociated protein	AGTRAP	57085	NP_065083	NM_020350	4D	1555736_a_at
CD209 antigen	CD209	30835	NP_066978	NM_021155	4D	1555729_a_at
regulator of Gprotein signalling 5	RGS5	8490	NP_003608	NM_003617	4D	1555725_a_at

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A DPribaculation factorille 2	ARI 3	403	NP 004302	NM 004311	4D	1568590 at
olfactory recentor family I subfamily I member	OR1J4	26219	NP 001004452	NM_001004452	4D	1567284_at
olfactory receptor, family 2, subfamily L, member	OR2L2	26246	NP_001004686	NM_001004686	4D	1567238_at
tropomyosin 4	TPM4	7171	NP_003281	NM_003290	4D	1567107_s_at
AT rich interactive domain 1B (SWIIlike)	ARIDIB	57492	NP_065783	NM_020732	4D	1566991_at
olfactory receptor, family 2, subfamily M, member	OR2M4	26245	NP_059974	NM_017504	4D	1566289_at
myoneurin	MYNN	55892	NP_061127	NM_018657	4D	1566109_at
dual oxidase 1	DUOXI	53905	NP_059130	NM_017434	4D	1565795_at
hypothetical protein LOC164045	LOC164045	164045	XP_371267	XM_371267	4D	1564285_at
chromosome 5 open reading frame 16	C5orf16	285613	NP_776189	NM_173828	4D	1564031_a_at
hypothetical protein FLJ32110	FLJ32110	219578	NP_857597	NM_181646	4D	1564022_at
plexin D1	PLXNDI	23129	NP_055918	NM_015103	4D	1563657_at
parvin, alpha	PARVA	55742	NP_060692	NM_018222	4D	1563458_at
hypothetical gene supported by BC040875	LOC400456	400456	XP_378545	XM_378545	4D	1562901_at
sterile alpha motif domain containing 12	SAMD12	401474	NP_997389	NM_207506	4D	1562637_at
hypothetical gene supported by AK097381; BC040866	LOC400680	400680	XP_378783	XM_378783	4D	1562348_at
CUB and Sushi multiple domains 1	CSMD1	64478	NP_150094	NM_033225	4D	1562329_at
pyruvate dehydrogenase kinase, isoenzyme 4	PDK4	5166	NP_002603	NM_002612	4D	1562321_at
chromosome 3 open reading frame 15	C3orf15	89876	NP_203528	NM_033364	4D	1561429 a at
hypothetical gene supported by BC042100	LOC440602	440602	XP_498751	XM_498751	4D	1561335_at
stereocilin	STRC	161497	NP_714544	NM_153700	4D	1561306_s_at
chromosome 21 open reading frame 106	C21orf106	23181	NP_996773	11	4D	1561286_a_at
hypothetical protein LOC338579	LOC338579	338579	XP_495807	XM_495807	4D	1561225_at
nitric oxide synthase 1 (neuronal)	NOSI	4842	NP_000611	NM_000620	4D	1560974_s_at
hypothetical protein LOC285972	LOC285972	285972	XP_379530	XM_379530	4D	1560762_at
Rhoguanine nucleotide exchange factor	RGNEF	64283	XP_371755	XM_371755	4D	1560348_at
laminin, alpha 3	LAMA3	3909	NP_937762	NM_198129	4D	1560078_at
Rho guanine nucleotide exchange factor (GEF) 12	ARHGEF12	23365	NP_056128	NM_015313	4D	1559833_at
hypothetical protein FLJ35681	FLJ35681	283897	NP_787096	NM_175900	4D	1559584 a at
KIAA1467 protein	KIAA1467	57613	XP_370682	XM_370682	4D	1559302_at
spergenl	SPAS1	219938	NP_777587	NM_174927	4D	1559138_a_at
KIAA1447 protein	KIAA1447	57597	XP_371084	XM_371084	4D	1559075_s_at
HLA complex group 27	HCG27	253018	NP_859068	NM_181717	4D	1559050_at
KIAA1240 protein	KIAA1240	54454	XP_039676	XM_039676	4D	1558807_at

ribosomal protein L28	KPL28	8010	NF_000982	166000 ININI	#5	200000 3 41
ribosomai protein L35	Nr Loo		MI 000000	- 1		
a) pomental protein (1.31313)	nni oz		ND 000140		4D	200002 at
hypothetical protein FI 113197	FLJ13197	79667	NP 078890	NM 024614	4D	1570414_x_at
hypothetical protein FLJ31413	FLJ31413	155061	NP_689770	NM_152557	4D	1570373_at
Ras homolog enriched in brain like 1	RHEBLI	121268	NP_653194	NM_144593	4D	1570253_a_at
keratin 6L	KRT6L	338785	NP_787028	NM_175834	4D	1569909_at
hypothetical gene supported by BC039326	LOC400605	400605	XP_378685	XM_378685	4D	1569805_at
nicotinamide nucleotide adenylyltransferase	NMNAT1	64802	NP_073624	NM_022787	4D	1569631_at
	FNDC3B	64778	NP_073600	NM_022763	4D	1569490_at
eukaryotic translation initiation factor 2C. 4	EIF2C4	192670	NP_060099	NM_017629	4D	1569408_at
chromosome 11 open reading frame 30	C11orf30	56946	NP_064578	NM_020193	4D	1569349_at
protein tyrosine phosphatase, receptor type. G	PTPRG	5793	NP_002832	NM_002841	4D	1569323_at
KIAA1731 protein	KIAA1731	85459	XP_374922	XM_374922	4D	1569302_at
zinc finger protein 505	ZNF505	81931	NP_112495	NM_031218	4D	1569240_at
hypothetical protein MGC40368	MGC40368	255394	NP_689985	NM_152772	4D	1569207_s_at
hypothetical protein MGC40368	MGC40368	255394	NP_689985	NM_152772	4D	1569206_at
hypothetical protein MGC29649	MGC29649	283237	NP_776171	NM_173810	4D	1569189_at
ATPbinding cassette, subfamily A (ABC1), member 4	ABCA4	24	NP_000341	NM_000350	40	1369102_at
dependen						15/0103
SWI/SNF related, matrix associated, actin	SMARCA4	6597	NP_003063	NM_003072	4D	1569073_x_at
IQ motif containing GTPase activating protein 3	IQGAP3	128239	NP_839943	NM_178229	4D	1569062_s_at
PRO0149 protein	PRO0149	29035	NP_054836	NM_014117	4D	1568954_s_at
hypothetical protein FLJ35834	FLJ35834	154865	NP_849149	NM_178827	4D	1568924_a_at
neighbor of BRCA1 gene 1	NBR1	4077	NP_114068	NM_031862	4D	1568856_at
zinc finger protein 505	ZNF505	81931	NP_112495	NM_031218	4D	1568720_at
TBC1 (tre2/USP6, BUB2, cdc16) domain family, membe	TBCIDI	23216	NP_055988	NM_015173	4D	1568/13_a_at
FGFR1 oncogene partner	FGFRIOP	111116	NP_008976	NM_007045	4D	1568678 s_at
similar to RIKEN cDNA 2310076L09	LOC440503	440503	XP_496282	XM_496282	4D	1568667_s_at
hypothetical gene supported by AK075484; BC014578	LOC339804	339804	XP_291016	XM_291016	4D	1568658_at
phosphoinositide3kinase, regulatory subunit 2 (p85	PIK3R2	5296	NP_005018	NM_005027	40	1568629_s_at
FLJ39739 protein	FLJ39739	388685	NP_997283	NM_207400	4D	1568609_s_at
hypothetical gene supported by BC031266	LOC400368	400368	XP_378511	XM_378511	4D	1568592_at

spectrin, beta, nonerythrocytic 1	SPIBNI	0/11	INF_UUSI 19	02120		8:
homolog,			NID COST IO	NIA 002120	È	200672 v at
ubiquitinconjugating enzyme E2D 3 (UBC4/5	UBE2D3	7323	155500_AN	NM_003340	4D	200669_s_at
translocase of outer mitochondrial membrane 20 hom	TOMM20	9804	NP_055580	NM_014765	41)	200662_s_at
signal sequence receptor, beta (transloconassociat	SSR2	6746	NP_003136	NM_003145		200652_at
glutamateammonia ligase (glutamine synthase)	GLUL	2752	NP_002056	NM_002065		
eukaryotic translation initiation factor 3, subuni	EIF3S8	8663	NP_003743	NM_003752		200647_x_at
superoxide dismutase 1, soluble (amyotrophic later	SODI	6647	NP_000445	NM_000454		200642_at
WD repeat domain 1	WDR1	9948	NP_059830	NM_017491		200609_s_at
RAD21 homolog (S. pombe)	RAD21	5885	NP_006256	NM_006265		200607_s_at
ribosomal protein L37	RPL37	6167	NP_000988	NM_000997	4D	200092 s_at
ribosomal protein S25	RPS25	6230	NP_001019	NM_001028	4D	200091_s_at
ribosomal protein S7	RPS7	6201	NP_001002	NM_001011	4D	200082_s_at
IK cytokine, downregulator of HLA II	IK	3550	NP_006074	NM_006083	4D	200066_at
ribosomal protein L30	RPL30	6156	NP_000980	NM_000989		200062_s_at
ras homolog gene family, member A	RHOA	387	NP_001655	NM_001664	4D	200059_s_at
nuclear DNAbinding protein	CID	10438	NP_006324	NM_006333	4D	200056_s_at
sperm associated antigen 7	SPAG7	9552	NP_004881	NM_004890	4D	200053_at
zinc finger protein 146	ZNF146	7705	NP_009076	NM_007145	4D	200050_at
defender against cell death 1	DADI	1603	NP_001335	NM_001344	40	200046_at
ATPbinding cassette, subfamily F (GCN20), member 1	ABCF1	23	NP_001081	NM_001090	4D	200045_at
enhancer of rudimentary homolog (Drosophila)	ERH	2079	NP_004441	NM_004450	40	200043_at
proteasome (prosome, macropain) subunit, beta type	PSMB2	5690	NP_002785	NM_002794	40	200039_s_at
ribosomal protein S11	RPS11	6205	NP_001006	NM_001015	4D	200031_s_at
ribosomal protein L19	RPL19	6143	NP_000972	NM_000981	4D	200029_at
ribosomal protein L34	RPL34	6164	NP_000986	NM_000995	4D	200026_at
ribosomal protein L27	RPL27	6155	NP_000979	NM_000988	4D	200025_s_at
ribosomal protein L18	RPL18	6141	NP_000970	NM_000979	4D	200022_at
ribosomal protein S13	RPS13	6207	NP_001008	NM_001017	4D	200018_at
ribosomal protein L24	RPL24	6152	NP_000977	NM_000986	4D	200013_at
ADPribosylation factor 3	ARF3	377	NP_001650	NM_001659	4D	200011_s_at
signal recognition particle 14kDa (homologous Alu	SRP14	6727	NP_003125	NM_003134	4D	200007_at

procollagenlysine 1, 20xoglutarate 5dioxygenase 1	PLODI	5351	NP_000293	NM_000302	4D	200827_at
small nuclear ribonucleoprotein D2 polypeptide 16.	SNRPD2	6633	NP_004588	NM_004597	4D	200826_at
ribosomal protein L29	RPL29	6159	NP_000983	NM_000992	4D	200823_x_at
ribosomal protein S15	RPS15	6209	NP_001009	NM_001018	4D	200819_s_at
ATP synthase, H+ transporting, mitochondrial F1 co	ATP5O	339	NP_001088	NM_00109/	ŧ	200010_41
ribosomal protein S10	RPS10	6204	NP_001005	NM 001014	3 5	18_X_/10007
heat shock 70kDa protein 1A	HSPAIA	3303	NP_005336	NM_005345	3 6	e s
heat shock 70kDa protein 1A	HSPAIA	3303	NP_005336	NM_005345	40	
myeloid cell leukemia sequence 1 (BCL2related)	MCL1	4170	NP_068779	NM_021960	4D	200798_x_at
myeloid cell leukemia sequence 1 (BCL2related)	MCL1	4170	NP_068779	NM_021960	40	200797_s_at
myeloid cell leukemia sequence 1 (BCL2related)	MCL1	4170	NP_068779	NM_021960	4D	200796_s_at
phosphoprotein enriched in astrocytes 15	PEA15	8682	NP_003759	NM_003768	4D	200788_s_at
ribosomal protein S15a	RPS15A	6210	NP_001010	NM_001019	4D	200781_s_at
methionine adenosyltransferase II, alpha	MAT2A	4144	NP_005902	NM_005911	4D	200768_s_at
ribosomal protein, large, P1	RPLP1	6176	NP_000994	NM_001003	4D	200763_s_at
calumenin	CALU	813	NP_001210	NM_001219	4D	200755_s_at
ferritin, heavy polypeptide I	FTHI	2495	NP_002023	NM_002032	4D	200748_s_at
guanine nucleotide binding protein (G protein), be	GNB1	2782	NP_002065	NM_002074	4D	200744_s_at
ribosomal protein S27 (metallopanstimulin 1)	RPS27	6232	NP_001021	NM_001030	4D	200741_s_at
nascentpolypeptideassociated complex alpha polypep	NACA	4666	NP_005585	NM_005594	4D	200735_x_at
ADPribosylation factor 3	ARF3	377	NP_001650	NM_001659	4D	200734_s_at
protein tyrosine phosphatase type IVA, member 1	PTP4A1	7803	NP_003454	NM_003463	4D	200733_s_at
ARP2 actinrelated protein 2 homolog (yeast)	ACTR2	10097	NP_005713	NM_005722	4D	200728_at
ribosomal protein L10	RPL10	6134	NP_006004	NM_006013	4D	200725_x_at
ribosomal protein L13a	RPL13A	23521	NP_036555	NM_012423	4D	200716_x_at
glutamicoxaloacetic transaminase 2, mitochondrial	GOT2	2806	NP_002071	NM_002080	4D	200708_at
dynein, cytoplasmic, light polypeptide 1	DNCLI	8655	NP_003737	NM_003746	4D	200703_at
NiemannPick disease, type C2	NPC2	10577	NP_006423	NM_006432	4D	200701_at
KDEL (LysAspGluLeu) endoplasmic reticulum protein	KDELR2	11014	NP_006845	NW_006854	45	200/00_s_at
gelsolin (amyloidosis, Finnish type)	GSN	2934	NP_000168	NM_000177	4D	200696_s_at
ribosomal protein L32	RPL32	6161	NP_000985	NM_000994	4D	200674_s_at

signal sequence receptor, delta (transloconassocia	SSR4	6748	NP_006271	NM_006280	40	201004_at
alanyltRNA synthetase	AARS	16		NM_001605	\$ 2	201000_at
cytoskeletonassociated protein 4	CKAP4	10970		NM_006825	145	201000 s_at
cytoskeletonassociated protein 4	CKAP4	10970	NP_006816	NM_006825	4D	200998_s_at
hypoxiainducible factor 1, alpha subunit (basic he	HIFIA	3091	NP_001521	NM_001530	40	200989_at
CD59 antigen p1820 (antigen identified by monoclon	CD59	966	NP_000602	NM_000611	#	
pyruvate dehydrogenase (lipoamide) alpha 1	PDHAI	5160	NP_000275	NM_000284	18	200980 s at
transmembrane 4 superfamily member 8	TM4SF8	10099	NP_005715	NM_005724	4D	S
transmembrane 4 superfamily member 8	TM4SF8	10099	NP_944492	NM_198902	4D	12
actin binding LIM protein 1	ABLIMI	3983	NP_006711	NM_006720	4D	200965_s_at
ribosomal protein L31	RPL31	6160	NP_000984	NM_000993	4D	200963_x_at
clathrin, light polypeptide (Lca)	CLTA	1211	NP_009027	NM_007096	4D	200960_x_at
fusion (involved in t(12;16) in malignant liposarc	FUS	2521	NP_004951	NM_004960	4D	200959_at
syndecan binding protein (syntenin)	SDCBP	6386	NP_005616	NM_005625	4D	200958_s_at
ribosomal protein S20	RPS20	6224	NP_001014	NM_001023	4D	200949_x_at
arginineglutamic acid dipeptide (RE) repeats	RERE	473	NP_036234	NM_012102	4D	200939_s_at
ribosomal protein S23	RPS23	6228	NP_001016	NM_001025	4D	200926_at
cytochrome c oxidase subunit VIa polypeptide 1	COX6A1	1337	NP_004364	NM_004373	4D	200925_at
solute carrier family 3 (activators of dibasic and	SLC3A2	6520	NP_002385	NM_002394	4D	200924_s_at
polyhomeoticlike 2 (Drosophila)	PHC2	1912	NP_004418	NM_004427	4D	200919_at
transgelin 2	TAGLN2	8407	NP_003555	NM_003564	4D	200916_at
protein phosphatase 1G (formerly 2C), magnesiumdep	PPM1G	5496	NP_002698	NM_002707	4D	200913_at
ribosomal protein, large P2	RPLP2	6181	NP_000995	NM_001004	4D	200909_s_at
major histocompatibility complex, class I, E	HLAE	3133	NP_005507	NM_005516	4D	200905_x_at
splicing factor, arginine/serinerich 10 (transform	SFRS10	6434	NP_004584	NM_004593	4D	200893_at
ribosomal protein L23	RPL23	9349	NP_000969	NM_000978	4D	200888_s_at
creatine kinase, brain	CKB	11152	NP_001814	NM_001823	4D	200884_at
nucleolar protein 5A (56kDa with KKE/D repeat)	NOL5A	10528	NP_006383	NM_006392	4D	200874 s_at
RABIIA, member RAS oncogene family	RABIIA	8766	NP_004654	NM_004663	4D	200863_s_at
nuclear receptor corepressor	NCOR1	9611	NP_006302	NM_006311	4D	200857_s_at
guanine nucleotide binding protein (C protein) he	GNB2	2783	NP_005264	NM_005273	4D	200852_x_at
glutamylprolyltRNA synthetase	EPRS	2058	NP_004437	NM_004446	4D	200843_s_at
ribosomal protein \$21	RPS21	6227	NP_001015	NM_001024	4D	200834_s_at

subcomplex,				**************************************		***************************************
NADH dahydroganogo (phiguingan) 1 ha	NDI IFR8	4714	NP 004995	NM 005004	4D	201227 s at
NADH dehydrogenase (ubiquinone) 1 beta subcomplex,	NDUFB8	4714	NP_004995	NM_005004	4D	201226_at
cystatin B (stefin B)	CSTB	1476	NP_000091	NM_000100	4D	201201_at
phosphatidylinositol transfer protein, alpha	PITPNA	5306	NP_006215	NM_006224	4D	201192_s_at
protease, serine, 11 (IGF binding)	PRSS11	5654	NP_002766	NM_002775	4D	201185_at
guanine nucleotide binding protein (G protein), al	GNAI3	2773	NP_006487	NM_006496	4D	201181_at
pumilio homolog 1 (Drosophila)	PUM1	9698	NP_055491	NM_014676	4D	201164 s at
Nmyristoyltransferase 1	NMTI	4836	NP_066565	NM_021079	4D	201158_at
RAB5C, member RAS oncogene family	RAB5C	5878	NP_004574	NM_004583	4D	201156_s_at
nuclear factor (erythroidderived 2)like 2	NFE2L2	4780	NP_006155	NM_006164	4D	201146_at
HS1 binding protein	HAXI	10456	NP_006109	NM_006118	4D	201145_at
major histocompatibility complex, class II, DP bet	HLADPB1	3115	NP_002112	NM_002121	4D	201137_s_at
cytochrome c oxidase subunit VIIc	COX7C	1350	NP_001858	NM_001867	4D	201134_x_at
heterogeneous nuclear ribonucleoprotein H2 (H')	HNRPH2	3188	NP_062543	NM_019597	4D	201132_at
splicing factor, arginine/serinerich 7, 35kDa	SFRS7	6432	NP_006267	NM_006276	4D	201129_at
cytochrome c oxidase subunit 8A (ubiquitous)	COX8A	1351	NP_004065	NM_004074	4D	201119_s_at
carboxypeptidase E	CPE	1363	NP_001864	NM_001873	4D	201117_s_at
CSE1 chromosome segregation 11ike (yeast)	CSEIL	1434	NP_803185	NM_177436	4D	2011111_at
thrombospondin 1	THBS1	7057	NP_003237	NM_003246	4D	201109_s_at
ribosomal protein S29	RPS29	6235	NP_001023	NM_001032	 4 D	201094_at
dynactin 1 (p150, glued homolog, Drosophila)	DCTNI	1639	NP_004073	NM_004082	4D	201082_s_at
general transcription factor II, i	GTF2I	2969	NP_001509	NM_001518	4D	201065_s_at
poly(A) binding protein, cytoplasmic 4 (inducible	PABPC4	8761	NP_003810	NM_003819	4D	201064 s at
heterogeneous nuclear ribonucleoprotein A0	HNRPA0	10949	NP_006796	NM_006805	4D	201054_at
dual specificity phosphatase 1	DUSP1	1843	NP_004408	NM_004417	4D	201044_x_at
guanine nucleotide binding protein (G protein), al	GNAI2	2771	NP_002061	NM_002070	4D	201040_at
L3hydroxyacylCoenzyme A dehydrogenase, short chain	HADHSC	3033	NP_005318	NM_005327	4D	201036_s_at
eukaryotic translation initiation factor 5B	EIF5B	9669	NP_056988	NM_015904	4D	201024_x_at
destrin (actin depolymerizing factor)	DSTN	11034	NP_006861	NM_006870	4D	201022 s_at
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM_006472	4D	201010_s_at
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM_006472	4D	201009_s_at
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM_006472	4D	201008_s_at

hydroxysteroid (17beta) dehydrogenase 4	HSD17B4	3295	NP_000405	NM_000414	140	201412_at
low density lipoprotein receptorrelated protein 10	LW 10	20020	TAL 000 105	LIVOU MIL		
	nida i	00000	NP 054764	NM 014045	4D	201412 at
ribosomal protein L36a	RPL36A	6173	NP_066357	NM_021029	4D	201406_at
adrenergic, beta, receptor kinase 1	ADRBK1	156	NP_001610	NM_001619	4D	201401_s_at
proteasome (prosome, macropain) subunit, beta type	PSMB3	5691	NP_002786	NM_002795	40	201400_at
insulinlike growth factor 2 receptor	IGF2R	3482	NP_000867	NM_000876	4D	201392_s_at
neighbor of BRCA1 gene 1	NBRI	4077	NP_005890	NM_005899	4D	201384_s_at
ubiquitin associated protein 2like	UBAP2L	9898	NP_055662	NM_014847	4D	201377_at
ornithine decarboxylase antizyme 2	OAZ2	4947	NP_002528	NM_002537	4D	201365_at
influenza virus NS1A binding protein	IVNSIABP	10625	NP_006460	NM_006469	4D	201363_s_at
coatomer protein complex, subunit beta	СОРВ	1315	NP_057535	NM_016451	4D	201359_at
coatomer protein complex, subunit beta	СОРВ	1315	NP_057535	NM_016451	4D	201358_s_at
YME1like I (S. cerevisiae)	YMEILI	10730	NP_055078	NM_014263	4D	201352_at
glutathione peroxidase 3 (plasma)	GPX3	2878	NP_002075	NM_002084	4D	201348_at
adiponectin receptor 2	ADIPOR2	79602	NP_078827	NM_024551	45	201346_at
ubiquitinconjugating enzyme E2D 2 (UBC4/5 homolog,	UBE2D2	7322	NP_003330	NM_003339	4D	201343_at
small nuclear ribonucleoprotein polypeptide C	SNRPC	6631	NP_003084	NM_003093	4D	201342_at
ectodermalneural cortex (with BTBlike domain)	ENCI	8507	NP_003624	NM_003633	4D	201341_at
vets erythroblastosis virus E26 oncogene homolog	ETS2	2114	NP_005230	NM_005239	4D	201328_at
EBNA1 binding protein 2	EBNA1BP2	10969	NP_006815	NM_006824	4D	201323_at
serine/threonine kinase 25 (STE20 homolog, yeast)	S1K25	10494	NF_006365	NM_0065/4	=	201314_at
chromosome 5 open reading frame 13	C5orf13	9315	NP_004763	NM_004772	3 5	201310_s_at
DEAD (AspGluAlaAsp) box polypeptide 48	DDX48	9775	NP_055555	NM_014740	40	201303_at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	4D	201293_x_at
proteasome (prosome, macropain) subunit, alpha typ	PSMA5	5686	NP_002781	NM_002790	4D	2012/4_at
aldoketo reductase family 1, member B1 (aldose red	AKRIBI	231	NP_001619	NM_001628	45	2012/2_at
ATPase, Na+/K+ transporting, beta 1 polypeptide	ATPIBI	481	NP_001668	NM_001677	4D	201242_s_at
capping protein (actin filament) muscle Zline, alp	CAPZA2	830	NP_006127	NM_006136	4D	201238_s_at
capping protein (actin filament) muscle Zline, alp	CAPZA2	830	NP_006127	NM_006136	4D	201237_at
BTG family, member 2	BTG2	7832	NP_006754	NM_006763	4D	201235_s_at

likely ortholog of mouse immediate early	LEKEPO4	22824	NP_060941	NIVI 0104/1	410	201070 s_at
eukaryotic translation initiation factor 3, subuni	EIFOOO	0007		NIM 010471		ء ا
Sprients racio promis gradunic ren (polybyrini	ElE3C3	7338	ND 0037/7	NM 003756	4D	201592 at
splicing factor proline/olutamine rich (polumurimi	SFPO	6421	NP 005057	NM 005066	4D	201585_s_at
Sec23 homolog B (S. cerevisiae)	SEC23B	10483	NP_006354	NM_006363	4D	201582_at
hypothetical protein DJ971N18.2	DJ971N18.2	56255	NP_066979	NM_021156	4D	201580_s_at
nonmetastatic cells 1, protein (NM23A) expressed i	NMEI	4830	NP_000260	NM_000269	4D	201577_at
dCMP deaminase	DCTD	1635	NP_001912	NM_001921	4D	201572_x_at
low molecular mass ubiquinonebinding protein (9.5k	QPC	27089	NP_055217	NM_014402	4D	201568_at
dual specificity phosphatase 3 (vaccinia virus pho	DUSP3	1845	NP_004081	NM_004090	4D	201538_s_at
ubiquitinlike 3	UBL3	5412	NP_009037	NM_007106	4D	201535_at
proteasome (prosome, macropain) subunit, alpha typ	PSMA3	5684	NP_002779	NM_002788	4D	201532_at
ATPase, H+ transporting, lysosomal 14kDa, V1 subun	ATP6V1F	9296	NP_004222	NM_004231	4D	201527_at
ubiquitinconjugating enzyme E2N (UBC13 homolog, ye	UBE2N	7334	NP_003339	NM_003348	4D	201524_x_at
small nuclear ribonucleoprotein polypeptide N	SNRPN	6638	NP_003088	NM_003097	4D	201522_x_at
translocase of outer mitochondrial membrane 70 hom	TOMM70A	9868	NP_055635	NM_014820	4D	201519_at
nuclear cap binding protein subunit 2, 20kDa	NCBP2	22916	NP_031388	NM_007362	4D	201517_at
protein phosphatase 1, regulatory (inhibitor) subu	PPP1R11	6992	NP_068778	NM_021959	4D	201500_s_at
ribosomal protein L41	RPL41	6171	NP_066927	NM_021104	4D	201492_s_at
AHA1, activator of heat shock 90kDa protein ATPase	AHSAI	10598	NP_036243	NM_012111	40	201491_at
peptidylprolyl isomerase F (cyclophilin F)	PPIF	10105	NP_005720	NM_005729	4D	201490_s_at
quiescin Q6	QSCN6	5768	NP_002817	NM_002826	4D	201482_at
sequestosome 1	SQSTM1	8878	NP_003891	NM_003900	4D	201471 s at
NAD(P)H dehydrogenase, quinone 1	NQ01	1728	NP_000894	NM_000903	4D	201468_s_at
Ras homolog enriched in brain	RHEB	6009	NP_005605	NM_005614	4D	201453_x_at
dihydropyrimidinaselike 3	DPYSL3	1809	NP_001378	NM_001387	4D	201430 s_at
ribosomal protein L37a	RPL37A	6168	NP_000989	NM_000998	4D	201429_s_at
methylosome protein 50	MEP50	79084	NP_077007	NM_024102	4D	201421 s_at
SRY (sex determining region Y)box 4	SOX4	6659	NP_003098	NM_003107	4D	201417_at
SRY (sex determining region Y)box 4	SOX4	6659	NP_003098	NM_003107	4D	201416_at

VIAAAAA	K1V VU/07	0813	ND 022280	NM 014774	3	1201778 s at
secretory carrier membrane protein 3	SCAMP3	10067	NP_005689	NM_005698	40	2017/1_at
small nuclear ribonucleoprotein polypeptide A	SNRPA	6626	NP_004587	NM_004596	4D	11
cytochrome c oxidase subunit VIc	COX6C	1345	NP_004365	NM_004374	4D	201754_at
tumor protein p53 (LiFraumeni syndrome)	TP53	7157	NP_000537	NM_000546	40	201/46_at
NADH dehydrogenase (ubiquinone) FeS protein 3, 30k	NDUFS3	4722	NP_004542	NM_004551	; 4D	201740_at
membraneassociated ring finger (C3HC4) 6	MARCH6	10299	NP_005876	NM_005885	4D	201737_s_at
Nacetylg						
UDPNacetvlalphaDgalactosamine:nolvnentide	GALNT1	2589	NP_065207	NM_020474	4D	201723_s_at
erythrocyte membrane protein band 4.1 like 2	EPB41L2	2037	NP_001422	NM_001431	4D	201719_s_at
cyclin D3	CCND3	896	NP_001751	NM_001760	4D	201700_at
DNA (cytosine5)methyltransferase 1	DNMT1	1786	NP_001370	NM_001379	4D	201697_s_at
nucleoside phosphorylase	NP	4860	NP_000261	NM_000270	4D	201695_s_at
apoptosis inhibitor 5	API5	8539	NP_006586	NM_006595	4D	201687_s_at
chromosome 14 open reading frame 92	C14orf92	9878	NP_055643	NM_014828	4D	201684_s_at
peptidase (mitochondrial processing) beta	РМРСВ	9512	NP_004270	NM_004279	4D	201682_at
arsenate resistance protein ARS2	ARS2	51593	NP_056992	NM_015908	4D	201680_x_at
DC12 protein	DC12	56941	NP_064572	NM_020187	4D	201678 s_at
ribosomal protein S17	RPS17	6218	NP_001012	NM_001021	4D	201665_x_at
ADPribosylation factorlike 1	ARLI	400	NP_001168	NM_001177	4D	201658_at
interferon gamma receptor 2 (interferon gamma tran	IFNGR2	3460	NP_005525	NM_005534	4D	201642_at
bone marrow stromal cell antigen 2	BST2	684	NP_004326	NM_004335	4D	201641_at
acid phosphatase 1, soluble	ACP1	52	NP_004291	NM_004300	4D	201630_s_at
aspartyltRNA synthetase	DARS	1615	NP_001340	NM_001349	4D	201623_s_at
staphylococcal nuclease domain containing 1	SNDI	27044	NP_055205	NM_014390	4D	201622_at
aldehyde dehydrogenase 9 family, member A1	ALDH9A1	223	NP_000687	NM_000696	4D	201612_at
nuclear phosphoprotein similar to S. cerevisiae PW	PWPI	111137	NP_008993	NM_007062	4D	201606_s_at
repressor of estrogen receptor activity	REA	111331	NP_009204	NM_007273	40	201600_at
ornithine aminotransferase (gyrate atrophy)	OAT	4942	NP_000265	NM_000274	4D	201599_at
keratin 18	KRT18	3875	NP_000215	NM_000224	4D	201596_x_at
protein phosphatase 4, regulatory subunit 1	PPP4R1	9989	NP_005125	NM_005134	4D	201594_s_at
response,					j	20:20

putative translation initiation factor	SUII	10209	NP_005/92	108C00_MIN	#	ZUZUZI_X_at
LanC lantibiotic synthetase component Clike 1 (bac	LANCLI	10314	NP_006046	NM_00500	7 5	102020 = 11 CO CO C
(mitochondrial			NIN OOCO	NIM OOCOEE	٦	מבחבחב
acetylCoenzyme A acyltransferase 2	ACAA2	10449	NP_006102	NM_006111	4D	202003_s_at
Ras suppressor protein 1	RSU1	6251	NP_036557	NM_012425	4D	201980_s_at
				NM_021122	4D	201963_at
decay accelerating factor for complement (CD55, Cr	DAF	1604	NP_000565	NM_0005/4	40	201923_8_at
solute carrier family 20 (phosphate transporter),	SLC20A1	6574	NP_005406	NM_005415	5 6	201025 = ±
hypothetical protein FLJ10618	FLJ10618	55186	NP_060625		\$ B	201917 s_at
G1 to S phase transition 1	GSPT1	2935	NP_002085		40	s
FERM, RhoGEF (ARHGEF) and pleckstrin domain protei	FARPI	10160	NP_005757	NM_005766	4D	201910_at
CTD (carboxyterminal domain, RNA polymerase II, po	CTDSPL	10217	NP_005799	NM_005808	4D	201905_s_at
interleukin 13 receptor, alpha 1	IL13RA1	3597	NP_001551	NM_001560	4D	201888_s_at
interleukin 13 receptor, alpha 1	IL13RA1	3597	NP_001551	NM_001560	4D	201887_at
paraoxonase 2	PON2	5445	NP_000296	NM_000305	4D	201876_at
myelin protein zerolike 1	MPZL1	9019	NP_003944	NM_003953	4D	201874_at
ORF	LOC51035	51035	NP_056937	NM_015853	4D	201871_s_at
transducin (beta)like 1Xlinked	TBL1X	6907	NP_005638	NM_005647	4D	201868_s_at
leucine rich repeat (in FLII) interacting protein	LRRFIP1	9208	NP_004726	NM_004735	4D	201861_s_at
proteoglycan 1, secretory granule	PRG1	5552	NP_002718	NM_002727	4D	201859_at
cell division cycle 25B	CDC25B	994	NP_068659	NM_021873	4D	201853_s_at
neural precursor cell expressed, developmentally d	NEDD8	4738	NP_006147	NM_006156	4D	201840_at
tumorassociated calcium signal transducer 1	TACSTD1	4072	NP_002345	NM_002354	4D	201839 s_at
protein kinase, AMPactivated, beta 1 noncatalytic	PRKABI	5564	NP_006244	NM_006253	4D	201835_s_at
neuroepithelial cell transforming gene 1	NETI	10276	NP_005854	NM_005863	4D	201830_s_at
neuroepithelial cell transforming gene 1	NETI	10276	NP_005854	NM_005863	4D	201829_at
ring finger protein 14	RNF14	9604	NP_004281	NM_004290	4D	201824_at
endoglin (OslerRenduWeber syndrome 1)	ENG	2022	NP_000109	NM_000118	4D	201809_s_at
valyltRNA synthetase 2	VARS2	7407	NP_006286	NM_006295	4D	201797_s_at
lamin B receptor	LBR	3930	NP_002287	NM_002296	4D	201795_at
vrel reticuloendotheliosis viral oncogene homolog	RELA	5970	NP_068810	NM_021975	4D	201783_s_at
ring finger protein 13	RNF13	11342	NP_009213	NM_007282	4D	201780_s_at

stromal cell derived factor receptor 1	SDFR1	27020	NP_059429	NM_017455	4D	202228_s_at
KIAA0907 protein	KIAA0907	22889	NP_055764	NM_014949	4D	202220_at
LSM3 homolog, U6 small nuclear RNA associated (S.	LSM3	27258	NP_055278	NM_014463	40	202209_at
autocrine motility factor receptor	AMFR	267	NP_001135	NM_001144	1	202203_s_at
myotubularin related protein 3	MTMR3	8897	NP_066576	NM_021090	4D	202198_s_at
LIM domain kinase 2	LIMK2	3985	NP_005560	NM_005569	40	202193_at
protein phosphatase 2, regulatory subunit B (B56),	PPP2R5A	5525	NP_006234	NM_006243	4D	202187_s_at
TAF9 RNA polymerase II, TATA box binding protein (TAF9	6880	NP_003178	NM_00318/	40	202108_at
tubulin, beta 3	TUBB3	10381	NP_006077	NM_006086	100	202154_x_at
neural precursor cell expressed, developmentally d	NEDD9	4739	NP_006394	NM_006403	4D	202149_at
adenylosuccinate lyase	ADSL	158	NP_000017	NM_000026	4D	202144 s_at
zinc finger, MYND domain containing 11	ZMYNDH	10771	NP_006615	NM_006624	4D	202136_at
vabl Abelson murine leukemia viral oncogene homolo	ABLI	25	NP_005148	NMI_003137	#2	202125_8_81
von Willebrand factor	VWF	7450	NP_000543	NM_000552	12	202112_at
cytochrome c oxidase subunit VIIb	COX7B	1349		NM_001866	ŧ	202110_at
peptidase D	PEPD	5184	NP_000276	NM_000285	#	202118 at
bromodomain containing 4	BRD4	23476	NP_055114	NM_014299	15	202102 s at
vral simian leukemia viral oncogene homolog B (ras	RALB	5899	NP_002872	NM_002881	5 4	8
(ras						
vral simian leukemia viral oncogene homolog B	RALB	5899	NP_002872	NM_002881	4D	202100_at
HMT1 hnRNP methyltransferaselike 1 (S. cerevisiae)	HRMT1L1	3275	NP_001526	NM_001535	4D	202098_s_at
OGT(OGIcNAc transferase)interacting protein 106 KD	OIP106	22906	NP_055780	NM_014965	40	202080_s_at
NADH dehydrogenase (ubiquinone) 1, alpha/beta subc	NDUFABI	4706	NP_004994	NM_005003	40	2020//_at
low density lipoprotein receptor (familial hyperch	LDLR	3949	NP_000518	NM_000527	41)	202068_s_at
sell suppressor of lin12like (C. elegans)	SELIL	6400	NP_005056	NM_005065	4D	202064_s_at
sell suppressor of lin12like (C. elegans)	SELIL	6400	NP_005056	NM_005065	4D	202061_s_at
TGFB1 induced antiapoptotic factor	TIAF1	9220	NP_004731	NM_004740	4D	202039_at
RB1inducible coiledcoil 1	RB1CC1	9821	NP_055596	NM_014781	4D	202034_x_at
ribosomal protein L38	RPL38	6169	NP_000990	NM_000999	4D	202029_x_at

retinoid X receptor, alpha	RXRA	6256	NP_002948	NM_002957	40	202426_S_at
acylCoA synthetase longchain family member 4	ACSL4	2182	NP_0/5266	NM_0229//	145	202427 S at
Yip1 interacting factor homolog (S. cerevisiae)	YIFI	10897		NM_020470	4D	at
(yeast)	FNFFSF	20141	N1_050777	1 1111 (1 (2 (2)		more france
cysteinyitkin A synthetase	CANS	26121	ND 056444	NM 015629	dD F	0
contain label - I decrease complex 3, signa 2 subunit	CADS	833	ND 001742	NM 001751	4D	202402 s at
adaptorrelated protein complex 3 sigma 2 subunit	AP3S2	10239	NP 005820	NM 005829	4D	202399_s_at
transcription elongation regulator	TCERG1	10915	NP_006697	NM_006706	4D	202396_at
ATPbinding cassette, subfamily F (GCN20), member 3	ABCF3	55324	NP_060828	NM_018358	4D	202394_s_at
phosphatidylserine decarboxylase	PISD	23761	NP_055153	NM_014338	4D	202392_s_at
leptin receptor overlapping transcript	LEPROT	54741	NP_059996	NM_017526	4D	202377_at
transcription elongation factor A (SII)like 4	TCEAL4	79921	NP_079139	NM_024863	4D	202371_at
torsin family 1, member A (torsin A)	TORIA	1861	NP_000104	NM_000113	4D	202349_at
fatty acid binding protein 5 (psoriasisassociated)	FABP5	2171	NP_001435	NM_001444	4D	202345_s_at
ubiquitinconjugating enzyme E2B (RAD6 homolog)	UBE2B	7320	NP_003328	NM_003337	4D	202335_s_at
ubiquitinconjugating enzyme E2B (RAD6 homolog)	UBE2B	7320	NP_003328	NM_003337	40	202334_s_at
polycystic kidney disease 1 (autosomal dominant)	PKD1	5310	NP_000287	NM_000296	† 0	202328_s_at
HLAB associated transcript 8	BAT8	10919	NP_006700	NM_006709	4D	202326_at
geranylgeranyl diphosphate synthase 1	GGPS1	9453	NP_004828	NM_004837	4D	202322 s_at
polymerase (RNA) II (DNA directed) polypeptide G	POLR2G	5436	NP_002687	NM_002696	4D	202306_at
NADH dehydrogenase (ubiquinone) I alpha subcomplex	NDUFA1	4694	NP_004532	NM_004541	40	202298_at
RER1 retention in endoplasmic reticulum 1 homolog	RERI	11079	NP_008964	NM_007033	#B	202296_s_at
laminin, gamma 2	LAMC2	3918	NP_005553	NM_005562	4D	202267_at
TRAF and TNF receptor associated protein	TTRAP	51567	NP_057698	NM_016614	4D	202266_at
hom						
translocase of outer mitochondrial mambrane 40	TOMM40	10452	NP 006105	NM 006114	4D	202264 s at
transcription factorlike	TCFLI	6944	NP_005988	NM_005997	4D	202261_at
phosphonoformate immunoassociated protein 5	PFAAP5	10443	NP_055702	NM_014887	4D	202259_s_at
cyclindependent kinase 4	CDK4	1019	NP_000066	NM_000075	4D	202246_s_at
nicotinamide Nmethyltransferase	TMNN	4837	NP_006160	NM_006169	4D	202237_at

DEAD (AspGluAlaAs) box polypeptide 19like	DDX19L	55308	NP_060802	NM_018332	40	2025//_s_at
small nuclear ribonucleoprotein D3 polypeptide 18k	SNRPD3	6634	NP_004166	NM_004175		a
supervillin	SVIL	6840	NP_003165	NM_003174		202565_s_at
DKFZP547E1010 protein	0 0	26097	NF_030422	NIM_01300/	;	202500 3 at
↓_	P29	20949	NF_050299	MM 015607		202560 s at
onydroxyomethylglutarylCoenzyme A reductase	HINIOCK	25040	ND 056200	NM 015484	4D	202553 s at
	HMGCB	3156	NP 000850	NM 000859	4D	202540 s at
	DKFZP5640123	25978	NP 054762	NM 014043	4D	202536_at
interferon regulatory factor 1	IRF1	3659	NP_002189	NM_002198	4D	202531_at
CCCTCbinding factor (zinc finger protein)	CTCF	10664	NP_006556	NM_006565	4D	202521_at
mutL homolog 1, colon cancer, nonpolyposis type	MLHI	4292	NP_000240	NM_000249	4D	202520_s_at
protein phosphatase 2, regulatory subunit B (B56),	PPP2R5D	5528	NP_006236	NM_006245	4D	202513_s_at
tumor necrosis factor, alphainduced protein 2	TNFAIP2	7127	NP_006282	NM_006291	4D	202509_s_at
acylCoenzyme A dehydrogenase, C4 to C12 straight c	ACADM	34	NP_000007	NM_000016	40	202302_at
solute carrier family 2 (facilitated glucose trans	SLC2A3	6515	NP_008862	NM_006931	40	202498_s_at
FXYD domain containing ion transport regulator 3	FXYD3	5349	NP_005962	NM_005971	4D	202488_s_at
RAN binding protein 1	RANBPI	5902	NP_002873	NM_002882	4D	202483_s_at
1 // 1				NM_006326	4D	202475_at
catenin (cadherinassociated protein), alphalike 1	CTNNALI	8727	NP_003789	NM_003798	4D	202468_s_at
6phosphofructo2kinase/fructose2,6biphosphatase	PFKFB3	5209	NP_004557	NM_004566	4D	202464_s_at
lipin 2	LPIN2	9663	NP_055461	NM_014646	4D	202459_s_at
chromosome 9 open reading frame 60	C9orf60	10444	NP_006327	NM_006336	4D	202456_s_at
retinoid X receptor, alpha	RXRA	6256	NP_002948	NM_002957	4D	202449_s_at
SPFH domain family, member 1	SPFH1	10613	NP_006450	NM_006459	4D	202444_s_at
adaptorrelated protein complex 3, sigma 1 subunit	AP3S1	1176	NP_001275	NM_001284	4D	202442_at
cytochrome P450, family 1, subfamily B, polypeptid	CYP1B1	1545	NP_000095	NM_000104	40	
vmyc myelocytomatosis viral oncogene homolog (avia	MYC	4609	NP_002458	NM_002467	4D	202431_S_at
diazepam binding inhibitor (GABA receptor modulato	DBI	1622	NP_065438	NM_020548	4D	202428_x_at
67 DKFZP564B167 protein	DKFZP564B167	25874	NP_056230	NM_015415	4D	202427_s_at

R3H domain (binds singlestranded nucleic acids)	R3HDM	23518	NP_056176	NM_015361	40	202/54_at
tryptophan rich basic protein	WRB	7485	NP_004618	NM_004627	4D	202749_at
guanylate binding protein 2, interferoninducible	GBP2	2634	NP_004111	NM_004120	4D	202748_at
integral membrane protein 2A	ITM2A	9452	NP_004858	NM_004867	4D	202747_s_at
latent transforming growth factor beta binding pro	LTBP1	4052	NP_000618	NM_000627	4D	202728_s_at
ligase I, DNA, ATPdependent	LIG1	3978	NP_000225	NM_000234	4D	202726_at
BET1 homolog (S. cerevisiae)	BET1	10282	NP_005859	NM_005868	4D	202710_at
cytochrome c oxidase subunit IV isoform 1	COX4II	1327	NP_001852	NM_001861	4D	202698_x_at
serine/threonine kinase 17a (apoptosisinducing)	STK17A	9263	NP_004751	NM_004760	4D	202695_s_at
upstream binding transcription factor, RNA polymer	UBTF	7343	NP_055048	NM_014233	4D	202692_s_at
small nuclear ribonucleoprotein D1 polypeptide 16k	SNRPD1	6632	NP_008869	NM_006938	4D	202690_s_at
general transcription factor IIA, 2, 12kDa	GTF2A2	2958	NP_004483	NM_004492	4D	202678_at
WiskottAldrich syndrome protein interacting protei	WASPIP	7456	NP_003378	NM_003387	4D	202665_s_at
WiskottAldrich syndrome protein interacting protei	WASPIP	7456	NP_003378	NM_003387	4D	202663_at
peroxisomal biogenesis factor 11B	PEXIIB	8799	NP_003837	NM_003846	4D	202658_at
membraneassociated ring finger (C3HC4) 7	MARCH7	64844	NP_073737	NM_022826	4D	202654_x_at
lysophosphatidylglycerol acyltransferase 1	LPGAT1	9926	NP_055688	NM_014873	4D	202651_at
ribosomal protein S19	RPS19	6223	NP_001013	NM_001022	4D	202649_x_at
tumor necrosis factor, alphainduced protein 3	TNFAIP3	7128	NP_006281	NM_006290	4D	202644_s_at
tumor necrosis factor, alphainduced protein 3	TNFAIP3	7128	NP_006281	NM_006290	4D	202643_s_at
ADPribosylation factorlike 3	ARL3	403	NP_004302	NM_004311	4D	202641_at
polymerase (RNA) II (DNA directed) polypeptide K,	POLR2K	5440	NP_005025	NM_005034	4D	202635_s_at
amyloid beta precursor protein (cytoplasmic tail)	APPBP2	10513	NP_006371	NM_006380	4D	202630_at
chromosome 14 open reading frame 11	C14orf11	55837	NP_060923	NM_018453	4D	202623_at
procollagenlysine, 20x0glutarate 5dioxygenase 2	PLOD2	5352	NP_000926	NM_000935	4D	202620_s_at
methyl CpG binding protein 2 (Rett syndrome)	MECP2	4204	NP_004983	NM_004992	4D	202618_s_at
cofactor required for Sp1 transcriptional activati	CRSP2	9282	NP_004220	NM_004229	4D	202611_s_at
nuclear receptor interacting protein 1	NRIP1	8204	NP_003480	NM_003489	4D	202600_s_at
membrane interacting protein of RGS16	MIR16	51573	NP_057725	NM_016641	4D	202593_s_at
singlestranded DNA binding protein 1	SSBP1	6742	NP_003134	NM_003143	4D	202591_s_at
heat shock 70kDa protein 1B	HSPAIB	3304	NP_005337	NM_005346	4D	202581_at

alanyl (membrane) aminopeptidase	ANPEP	290	NP_001141	NM_001150	4D	202888 s_at
DNAdamageinducible transcript 4	DDIT4	54541	NP_061931	NM_019058	4D	202887_s_at
nucleolar protein 7, 27kDa	NOL7	51406	NP_057251	NM_016167	4D	202882_x_at
DnaJ (Hsp40) homolog, subfamily B, member 12	DNAJB12	54788	NP_001002762	NM_001002762	4D	202866_at
transmembrane protein 4	TMEM4	10330	NP_055070	NM_014255	4D	202857_at
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP_004198	NM_004207	4D	202856_s_at
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP_004198	NM_004207	4D	202855_s_at
phosphatidylinositol glycan, class C	PIGC	5279	NP_002633	NM_002642	4D	202846_s_at
subcomplex,						
NADH dehydrogenase (ubiquinone) I beta	NDUFB7	4713	NP_004137	NM_004146	4D	202839_s_at
fucosidase, alphaL 1, tissue	FUCA1	2517	NP_000138	NM_000147	4D	202838_at
serine (or cysteine) proteinase inhibitor, clade A	SERPINAI	5265	NP_000286	NM_000295	4D	202833_s_at
GRIP and coiledcoil domain containing 2	GCC2	9648	NP_055450	NM_014635	4D	202832_at
LIM domain containing preferred translocation part	LPP	4026	NP_005569	NM_0055/8	4D	202822_at
transcription elongation factor B (SIII), polypept	TCEB3	6924	NP_003189	NM_003198	\$ D	202818_s_at
synovial sarcoma translocation, chromosome 18	SS18	6760	NP_005628	NM_005637	10	202817_s_at
TAR (HIV) RNA binding protein 1	TARBPI	6894	NP_005637	NM_005646	4D	202813_at
STAM binding protein	STAMBP	10617	NP_006454	NM_006463	4D	202811_at
developmentally regulated GTP binding protein 1	DRG1	4733	NP_004138	NM_004147	4D	202810_at
drebrin l	DBNI	1627	NP_004386	NM_004395	4D	202806_at
solute carrier family 1 (glial high affinity gluta	SLC1A3	6507	NP_004163	NM_004172	4D	202800_at
phospholipase C, gamma 1	PLCG1	5335	NP_877963	NM_182811	4D	202789_at
mitogenactivated protein kinaseactivated protein k	МАРКАРК3	7867	NP_004626	NM_004635	4D	202787_s_at
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFA7	4701	NP_004992	NM_005001	4D	202785_at
skeletal muscle and kidney enriched inositol phosp	SKIP	51763	NP_057616	NM_016532	4D	202781_s_at
soc2 suppressor of clear homolog (C. elegans)	SHOC2	8036	NP_031399	NM_007373	4D	202777_at
splicing factor, arginine/serinerich 8 (suppressor	SFRS8	6433	NP_004583	NM_004592	4D	202775_s_at
cyclin G2	CCNG2	901	NP_004345	NM_004354	4D	202770_s_at
acid phosphatase 2, lysosomal	ACP2	53	NP_001601	NM_001610	4D	202767_at
fibrillin 1 (Marfan syndrome)	FBN1	2200	NP_000129	NM_000138	4D	202766_s_at
PALM2AKAP2 protein	PALM2AKAP2	445815	NP_009134	NM_007203	4D	202759_s_at
regulatory factor Xassociated ankyrincontaining pr	RFXANK	8625	NP_003712	NM_003721	4D	202758_s_at
co						

breast carcinoma amplified sequence 2	BCAS2	10286	NP_005863	NM_005872	4D	203053_at
timeless homolog (Drosophila)	TIMELESS	8914	NP_003911	NM_003920	4D	203046_s_at
ninjurin 1	NINJI	4814	NP_004139	NM_004148	4D	203045_at
lysosomalassociated membrane protein 2	LAMP2	3920	NP_002285	NM_002294	4D	203042_at
uroporphyrinogen III synthase (congenital erythrop	UROS	7390	NP_000366	NM_000375	40	1
protein tyrosine phosphatase, receptor type, N pol	PTPRN2	5799	NP_002838	NM_002847	4D	203030_s_at
protein tyrosine phosphatase, receptor type, N pol	PTPRN2	5799	NP_002838	NM_002847	4D	203029_s_at
zinc finger and BTB domain containing 5	ZBTB5	9925	NP_055687	NM_014872	4D	203026_at
hypothetical protein HSPC111	HSPC111	51491	NP_057475	NM_016391	4D	203023_at
synovial sarcoma, X breakpoint 2 interacting prote	SSX2IP	117178	NP_054740	NM_014021	4D	203019_x_at
ribosomal protein L23a	RPL23A	6147	NP_000975	NM_000984	4D	203012_x_at
membrane protein, palmitoylated 1, 55kDa	MPPI	4354	NP_002427	NM_002436	4D	202974_at
dualspecificity tyrosine(Y)phosphorylation regulat	DYRK2	8445	NP_006473	NM_006482	4D	202971_s_at
dualspecificity tyrosine(Y)phosphorylation regulat	DYRK2	8445	NP_006473	NM_006482	4D	202968_s_at
calpain 6	CAPN6	827	NP_055104	NM_014289	4D	202966_at
regulatory factor X, 5 (influences HLA class II ex	RFX5	5993	NP_000440	NM_000449	4D	202963_at
A I P synthase, H+ transporting, mitochondrial F0 co	ATP3J2	9331	NF_004880	NN _004889	ŧ	202701_3_at
Nacetylgalactosaminidase, alpha	NAGA	4668	NP_000253	NM_000262	3 6	202945_S_at
homolog 1						
vyes1 Yamaguchi sarcoma viral oncogene	YES1	7525	NP_005424	NM_005433	4D	202933_s_at
neuroblastomaamplified protein	NAG	51594	NP_056993	NM_015909	4D	202926_at
glutamatecysteine ligase, catalytic subunit	GCLC	2729	NP_001489	NM_001498	4D	202922_at
ankyrin 2, neuronal	ANK2	287	NP_001139	NM_001148	4D	202921_s_at
CD97 antigen	CD97	976	NP_001775	NM_001784	4D	202910_s_at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP_002476	NM_002485	4D	202907_s_at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP_002476	NM_002485	4D	202906_s_at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP_002476	NM_002485	4D	202905_x_at
cathepsin S	CTSS	1520	NP_004070	NM_004079	4D	202902_s_at
				NM_004648	4D	202896_s_at
protein tyrosine phosphatase, nonreceptor type sub	PTPNSI	140885	NP_542970	NM_080792	4D	202895_s_at
CDC23 (cell division cycle 23, yeast, homolog)	CDC23	8697	NP_004652	NM_004661	4D	202892_at
microtubuleassociated protein 7	MAP7	9053	NP_003971	NM_003980	4D	202889_x_at
(aminopeptidase N						

syntaxin binding protein 3	STXBP3	6814	NP_009200	NM_007269	4D	203310_at
Jumonji, AT rich interactive domain 2	JARID2	3720	NP_004964	NM_004973	4D	203298 s_at
vacuolar protein sorting 11 (yeast)	VPS11	55823	NP_068375	NM_021729	4D	203292_s_at
DNA fragmentation factor, 45kDa, alpha polypeptide	DFFA	1676	NP_004392	NM_004401	4D	203277_at
chromosome 6 open reading frame 74	C6orf74	51020	NP_057147	NM_016063	4D	203259_s_at
Fc fragment of IgG binding protein	FCGBP	8857	NP_003881	NM_003890	4D	203240_at
sarcoma amplified sequence	SAS	6302	NP_005972	NM_005981	4D	203227_s_at
adenine phosphoribosyltransferase	APRT	353	NP_000476	NM_000485	4D	203219_s_at
HIV1 rev binding protein 2	HRB2	11103	NP_008974	NM_007043	4D	203202_at
5methyltetrahydrofolatehomocysteine methyltransfer	MTRR	4552	NP_002445	NM_002454	4D	203199_s_at
				NM_003138	4D	203182_s_at
glycine amidinotransferase (Larginine:glycine amid	GATM	2628	NP_001473	NM_001482	4D	203178_at
esophageal cancer associated protein	MGC16824	57020	NP_064710	NM_020314	4D	203173_s_at
KIAA0409 protein	KIAA0409	23378	NP_056139	NM_015324	4D	203171_s_at
cAMP responsive element binding proteinlike 1	CREBL1	1388	NP_004372	NM_004381	4D	203168_at
tissue inhibitor of metalloproteinase 2	TIMP2	7077	NP_003246	NM_003255	4D	203167_at
A kinase (PRKA) anchor protein 11	AKAPII	11215	NP_057332	NM_016248	4D	203156_at
SET domain, bifurcated 1	SETDB1	9869	NP_036564	NM_012432	4D	203155_at
KIAA0040 gene product	KIAA0040	9674	NP_055471	NM_014656	4D	203144_s_at
TATA box binding protein	ТВР	6908	NP_003185	NM_003194	4D	203135_at
Sec61 beta subunit	SEC61B	10952	NP_006799	NM_006808	4D	203133_at
inositol(myo)1(or 4)monophosphatase 2	IMPA2	3613	NP_055029	NM_014214	4D	203126_at
ubiquitin specific protease 52	USP52	9924	NP_055686	NM_014871	4D	203117_s_at
				NM_014247	4D	203097_s_at
mitochondrial translational initiation factor 2	MTIF2	4528	NP_002444	NM_002453	4D	203095_at
protease, serine, 25	PRSS25	27429	NP_037379	NM_013247	4D	203089_s_at
component of oligomeric golgi complex 2	COG2	22796	NP_031383	NM_007357	4D	203073_at
kelchlike 21 (Drosophila)	KLHL21	9903	NP_055666	NM_014851	4D	203068_at
				NM_014863	4D	203066_at
forkhead box K2	FOXK2	3607	NP_004505	NM_004514	4D	203064_s_at
3'phosphoadenosine 5'phosphosulfate synthase 2	PAPSS2	9060	NP_004661	NM_004670	4D	203060_s_at
3'phosphoadenosine 5'phosphosulfate synthase 2	PAPSS2	9060	NP_004661	NM_004670	4D	203058_s_at

IQ motif containing GTPase activating protein 2	IQGAP2	88/01	NP_006624	INIVI_UU0000	40	2024/4_41
solute carrier organic anion transporter family, m	SLCO2B1	11309	NE_007(31	0027700 MIN		ء [د
pleckstrin	PLEK	11200	NP_002033	120200 MIN		11
DICCKSITIN		5341	11 002000	122COU MIN	ة اد	
Cyclindependent sinase (CDC2118e) 10	DI EV	\$2/1		NM 002664	4D	را مر ا
cyclindenendent kingse (CDC21116) 10	CDK 10	8558	NP 003665	NM 003674	4D	203469 s at
presenilin 1 (Alzheimer disease 3)	PSEN1	5663	NP_015557	NM_007318	4D	203460_s_at
spermidine/spermine N1 acetyltransferase	SAT	6303	NP_002961	NM_002970	4D	203455_s_at
LIM domain binding 1	LDB1	8861	NP_003884	NM_003893	4D	203451_at
PKD2 interactor, golgi and endoplasmic reticulum a	PGEAI	25776	NP_056188	NM_015373	40	203450_at
cadherin 2, type 1, Ncadherin (neuronal)	CDH2	1000	NP_001783	NM_001792	4D	203440_at
ribonuclease P/MRP 30kDa subunit	RPP30	10556	NP_006404	NM_006413	4D	203436_at
membrane metalloendopeptidase (neutral endopeptida	MME	4311	NP_009218	NM_007287	4D	203435_s_at
family with sequence similarity 8, member A1	FAM8A1	51439	NP_057339	NM_016255	4D	203420_at
CD53 antigen	CD53	963	NP_000551	NM_000560	4D	203416_at
Down syndrome critical region gene 2	DSCR2	8624	NP_003711	NM_003720	4D	203405_at
transferrin	TF	7018	NP_001054	NM_001063	4D	203400_s_at
pregnancy specific betalglycoprotein 3	PSG3	5671	NP_066296	NM_021016	4D	203399 x_at
UDPNacetylalphaDgalactosamine:polypeptide Nacetylg	GALNT3	2591	NP_004473	NM_004482	4D	203397_s_at
kinesin family member 3C	KIF3C	3797	NP_002245	NM_002254	40	203390_s_at
kinesin family member 3C	KIF3C	3797	NP_002245	NM_002254	4D	203389_at
diacylglycerol kinase, alpha 80kDa	DGKA	1606	NP_001336	NM_001345	4D	203385_at
cell division cycle 40 homolog (yeast)	CDC40	51362	NP_056975	NM_015891	4D	203376_at
tripeptidyl peptidase II	TPP2	7174	NP_003282	NM_003291	4D	203375_s_at
tripeptidyl peptidase II	TPP2	7174	NP_003282	NM_003291	4D	203374_s_at
cmyc binding protein	MYCBP	26292	NP_036465	NM_012333	4D	203361_s_at
enhancer of zeste homolog 2 (Drosophila)	EZH2	2146	NP_004447	NM_004456	4D	203358_s_at
adaptorrelated protein complex 1, gamma 1 subunit	APIGI	164	NP_001119	NM_001128	4D	203350_at
caveolin 2	CAV2	858	NP_001224	NM_001233	4D	203324_s_at
zinc finger protein 148 (pHZ52)	ZNF148	7707	NP_068799	NM_021964	4D	203319_s_at
small nuclear ribonucleoprotein polypeptide E	SNRPE	6635	NP_003085	NM_003094	4D	203316_s_at
GTP binding protein 6 (putative)	GTPBP6	8225	NP_036359	NM_012227	4D	203314_at

coilin	COIL	8161	NP_004636	NM_004645	40	200000 S_at
protein C receptor, endothelial (EPCR)	PROCR	10544				at
Down syndrome critical region gene 3	DSCR3	10311	NP_006043	NM_006052		
carnitine palmitoyltransferase 1A (liver)	CPTIA	1374	NP_001867	NM_001876		203634 s_at
ELK1, member of ETS oncogene family	ELKI	2002	NP_005220	NM_005229		203617_x_at
polymerase (DNA directed), beta	POLB	5423	NP_002681	NM_002690	4D	203616_at
UTP14, U3 small nucleolar ribonucleoprotein, homol	UTP14C	9724	NP_067677	NM_021645	41)	205614_at
telomeric repeat binding factor 2	TERF2	7014	NP_005643	NM_005652		203611_at
inositol polyphosphate5phosphatase F	INPP5F	22876	NP_055752	NM_014937		203607_at
WW domain binding protein 4 (formin binding protei	WBP4	111193	NP_009118	NM_007187		203599_s_at
WW domain binding protein 4 (formin binding protei	WBP4	11193	NP_009118	NM_007187		203597_s_at
unc50 homolog (C. elegans)	UNC50	25972	NP_054763	NM_014044		203583_at
solute carrier family 7 (cationic amino acid trans	SLC7A6	9057	NP_003974	NM_003983	4D	203580_s_at
solute carrier family 7 (cationic amino acid trans	SLC7A6	9057	NP_003974	NM_003983	4D	203579_s_at
casein kinase 2, alpha prime polypeptide	CSNK2A2	1459	NP_001887	NM_001896	Đ	203575_at
nuclear factor, interleukin 3 regulated	NFIL3	4783	NP_005375	NM_005384	4D	203574_at
oralfacialdigital syndrome 1	OFD1	8481	NP_003602	NM_003611	4D	203569_s_at
Fanconi anemia, complementation group G	FANCG	2189	NP_004620	NM_004629	4D	203564_at
zinc fingers and homeoboxes 2	ZHX2	22882	NP_055758	NM_014943	4D	203556_at
CD4 antigen (p55)	CD4	920	NP_000607	NM_000616	4D	203547_at
calcium modulating ligand	CAMLG	819	NP_001736	NM_001745	4D	203538_at
LSMI homolog, U6 small nuclear RNA associated (S.	IWST	27257	NP_055277	NM_014462	4D	203534_at
adenomatosis polyposis coli	APC	324	NP_000029	NM_000038	4D	203527_s_at
phosphomevalonate kinase	PMVK	10654	NP_006547	NM_006556	4D	203515_s_at
tumor necrosis factor receptor superfamily, member	TNFRSF1B	7133	NP_001057	NM_001066	4D	203508_at
EPH receptor A2	EPHA2	1969	NP_004422	NM_004431	4D	203499_at
PPAR binding protein	PPARBP	5469	NP_004765	NM_004774	4D	203497_at
translokin	PIG8	9702	NP_055494	NM_014679	4D	203494_s_at
Sec61 gamma subunit	SEC61G	23480	NP_055117	NM_014302	4D	203484_at
NADH dehydrogenase (ubiquinone) 1, subcomplex unkn	NDUFC1	4717	NP_002485	NM_002494	4D	203478_at

small nuclear ribonucleoprotein polypeptide F	SNRPF	6636	NP_003086	NM_003095	40	203832_at
KIAA1002 protein	KIAA1002	22864	NP_055740	NM_014925	4D	203831_at
WD40 repeat protein Interacting with phospholnosit	WIPI49	55062	NP_060453	NM_017983	4D	203827_at
bromodomain containing 3	BRD3	8019	NP_031397	NM_007371	4D	203825_at
IGFII mRNAbinding protein 3	IMP3	10643	NP_006538	NM_006547	4D	203820_s_at
slit homolog 3 (Drosophila)	SLIT3	6586	NP_003053	NM_003062	4D	203813_s_at
DnaJ (Hsp40) homolog, subfamily B, member 4	DNAJB4	11080	NP_008965	NM_007034	4D	203810_at
NOL1/NOP2/Sun domain family, member 5	NSUN5	55695	NP_060514	NM_018044	4D	203802_x_at
mitochondrial ribosomal protein S14	MRPS14	63931	NP_071383	NM_022100	4D	203800_s_at
Bcell CLL/lymphoma 7A	BCL7A	605	NP_066273	NM_020993	4D	203795_s_at
mitochondrial ribosomal protein L33	MRPL33	9553	NP_004882	NM_004891	4D	203781_at
leiomodin I (smooth muscle)	LMODI	25802	NP_036266	NM_012134	4D	203766_s_at
grancalcin, EFhand calcium binding protein	GCA	25801	NP_036330	NM_012198	4D	203765_at
Srclikeadaptor	ALA	6503	NP_006739	NM_006748	4D	203761_at
Srclikeadaptor	SLA	6503	NP_006739	NM_006748	4D	203760_s_at
transcription factor 4	TCF4	6925	NP_003190	NM_003199	4D	203753_at
RNA binding motif, single stranded interacting pro	RBMS1	5937	NP_058523	NM_016839	40	203748_x_at
holocytochrome c synthase (cytochrome c hemelyase)	HCCS	3052	NP_005324	NM_00333	40	203/45_at
gamma	PPRC1	23082	NP_0558//	NM_010062	#	203/3/_s_at
KIAA0020	KIAA0020	9933	NP_055693	NM_014878	40	203712_at
phosphodiesterase 4B, cAMPspecific (phosphodiester	PDE4B	5142	NP_002591	NM_002600	4D	203708_at
hypothetical protein FLJ20244	FLJ20244	55621	NP_060192	NM_017722	4D	203701_s_at
tubulin, gamma complex associated protein 3	TUBGCP3	10426	NP_006313	NM_006322	4D	203690_at
transmembrane emp24 domain containing 1	TMEDI	81011	NP_006849	NM_006858	4D	203679_at
KIAA1018 protein	KIAA1018	22909	NP_055782	NM_014967	4D	203678_at
nucleobindin 2	NUCB2	4925	NP_005004	NM_005013	4D	203675_at
thiopurine Smethyltransferase	TPMT	7172	NP_000358	NM_000367	4D	203672_x_at
mannosidase, alpha, class 2C, member 1	MAN2CI	4123	NP_006706	NM_006715	4D	203668_at
tubulinspecific chaperone a	TBCA	6902	NP_004598	NM_004607	4D	203667_at
Xray repair complementing defective repair in Chin	XRCC1	7515	NP_006288	NM_006297	4D	203655_at

nucleotide binding protein 1 (MinD homolog, E.	NUBPI	4682	NP_002475	NM_002484	45	2039/8_at
CCAAT/enhancer binding protein (C/EBP), delta	CEBPD	1052	NP_005186	NM_005195	£	
CDC6 cell division cycle 6 homolog (S. cerevisiae)	CDC6	990	NP_001245	NM_001254	4D	
protein phosphatase 1A (formerly 2C), magnesiumdep	PPMIA	5494	NP_066283	NM_021003	4D	203966_s_at
Nmyc (and STAT) interactor	NMI	9111	NP_004679	NM_004688	4D	203964_at
calponin 1, basic, smooth muscle	CNNI	1264	NP_001290	NM_001299	4D	203951_at
butyrophilin, subfamily 2, member A1	BTN2A1	11120	NP_008980	NM_007049	4D	203944_x_at
kinesin family member 3B	KIF3B	9371	NP_004789	NM_004798	4D	203943_at
TATA box binding protein (TBP)associated factor, R	TAFIC	9013	NP_005670	NM_005679	4D	203938_s_at
matrix metalloproteinase 9 (gelatinase B, 92kDa ge	MMP9	4318	NP_004985	NM_004994	4D	203936_s_at
ATP synthase, H+ transporting, mitochondrial F1 co	ATP5D	513	NP_001678	NM_001687	4D	203926_x_at
carbohydrate (Nacetylglucosamine6O) sulfotransfera	CHST2	9435	NP_004258	NM_004267	4D	203921_at
transcription elongation factor A (SII), 2	TCEA2	6919	NP_003186	NM_003195	4D	203919_at
hydroxyprostaglandin dehydrogenase 15(NAD)	HPGD	3248	NP_000851	NM_000860	4D	203914_x_at
hydroxyprostaglandin dehydrogenase 15(NAD)	HPGD	3248	NP_000851	NM_000860	4D	203913_s_at
PTPL1associated RhoGAP 1	PARGI	9411	NP_004806	NM_004815	4D	203910_at
kangai I (suppression of tumorigenicity 6, prostat	KAII	3732	NP_002222	NM_002231	4D	203904_x_at
				NM_024547	4D	203900_at
TAF9 RNA polymerase II, TATA box binding protein (TAF9	6880	NP_057367	NM_016283	4D	203893_at
deathassociated protein kinase 3	DAPK3	1613	NP_001339	NM_001348	4D	203890_s_at
RAB21, member RAS oncogene family	RAB21	23011	NP_055814	NM_014999	4D	203885_at
COX17 homolog, cytochrome c oxidase assembly prote	COX17	10063	NP_005685	NM_005694	4D	203880_at
SUMO1/sentrin/SMT3 specific protease 3	SENP3	26168	NP_056485	NM_015670	4D	203871_at
vascular cell adhesion molecule 1	VCAM1	7412	NP_001069	NM_001078	4D	203868_s_at
for protein disulfide isomeraserelated	PDIR	10954	NP_006801	NM_006810	4D	203857_s_at
WD repeat domain 47	WDR47	22911	NP_055784	NM_014969	4D	203855_at
basic leucine zipper nuclear factor 1 (JEM1)	BLZF1	8548	NP_003657	NM_003666	4D	203840_at
glycoprotein A repetitions predominant	GARP	2615	NP_005503	NM_005512	4D	203835_at

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RNA binding motif protein 14	RBM14	10432	NP_006319	NM_006328	4D	204178_s_at
kinetochore associated 2	KNTC2	10403	NP_006092	NM_006101	4D	at
transcription factor Dp1	TFDPI	7027	NP_009042	NM_007111	4D	204147_s_at
enolase superfamily member 1	ENOSF1	55556	NP_059982	NM_017512	4D	204143_s_at
enolase superfamily member 1	ENOSFI	55556	NP_059982	NM_017512	4D	204142_at
tyrosylprotein sulfotransferase 1	TPST1	8460	NP_003587	NM_003596	4D	204140_at
forkhead box O3A	FOXO3A	2309	NP_001446	NM_001455	4D	204132_s_at
forkhead box O3A	FOXO3A	2309	NP_001446	NM_001455	4D	204131_s_at
replication factor C (activator 1) 3, 38kDa	RFC3	5983	NP_002906	NM_002915	4D	204128 s_at
replication factor C (activator 1) 3, 38kDa	RFC3	5983	NP_002906	NM_002915	4D	204127_at
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFAFI	51103	NP_057097	NM_016013	4D	204125_at
prolyl endopeptidase	PREP	5550	NP_002717	NM_002726	4D	204117_at
RNA binding motif protein, Xlinked 2	RBMX2	51634	NP_057108	NM_016024	4D	204098_at
RNA binding motif protein, Xlinked 2	RBMX2	51634	NP_057108	NM_016024	4D	204097_s_at
solute carrier family 5 (sodiumdependent vitamin t	SLC5A6	8884	NP_066918	NM_021095	4D	204087_s_at
glycine, glutamate, thienylcyclohexylpiperidinehin	KIAA0562	9731	NP_055519	NM_014704	4D	204075_s_at
topoisomerase I binding, arginine/serinerich	TOPORS	10210	NP_005793	NM_005802	4D	204071 s_at
Meis1, myeloid ecotropic viral integration site	MEIS1	4211	NP_002389	NM_002398	4D	204069_at
unc51like kinase 2 (C. elegans)	ULK2	9706	NP_055498	NM_014683	4D	204063 s_at
unc51like kinase 2 (C. elegans)	ULK2	9706	NP_055498	NM_014683	4D	204062_s_at
phosphatase and tensin homolog (mutated in multipl	PTEN	5728	NP_000305	NM_000314	45	12
phosphatase and tensin homolog (mutated in multipl	PTEN	5728	NP_000305	NM_000314	40	204053_x_at
WAS protein family, member 3	WASF3	10810	NP_006637	NM_006646	4D	204042_at
purinerich element binding protein A	PURA	5813	NP_005850	NM_005859	4D	204021_s_at
purinerich element binding protein A	PURA	5813	NP_005850	NM_005859	4D	204020_at
sprouty homolog 2 (Drosophila)	SPRY2	10253	NP_005833	NM_005842	4D	204011_at
PRKC, apoptosis, WT1, regulator	PAWR	5074	NP_002574	NM_002583	4D	204004_at
guanine nucleotide binding protein (G protein), be	GNB5	10681	NP_057278	NM_016194	4D	204000_at
ATPbinding cassette, subfamily D (ALD), member 4	ABCD4	5826	NP_064719	NM_020323	40	203981_s_at
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bromodomain and PHD finger containing, 1	BRPF1	7862	NP_004625	NM_004634	4D	204481_at
apolipoprotein Al	APOAI	335	NP_000030	NM_000039	40	204450_x_at
arachidonate 5lipoxygenase	ALOX5	240	NP_000689	NM_000698	4D	204445_s_at
PH domaincontaining protein	pp9099	80301	NP_079477	NM_025201	4D	204436_at
				NM_014778	4D	204435_at
muskelin 1, intracellular mediator containing kelc	MKLNI	4289	NP_037387	NM_013255	4D	204423_at
apolipoprotein CI	APOC1	341	NP_001636	NM_001645	4D	204416_x_at
neurofilament, heavy polypeptide 200kDa	NEFH	4744	NP_066554	NM_021076	4D	204412_s_at
echinoderm microtubule associated protein like 2	EML2	24139	NP_036287	NM_012155	4D	204398_s_at
G proteincoupled receptor kinase 5	GRK5	2869	NP_005299	NM_005308	4D	204396_s_at
mitochondrial ribosomal protein 63	MRP63	78988	NP_076931	NM_024026	4D	204387_x_at
fibroblast growth factor receptor 3 (achondroplas)	FGFR3	2261	NP_000133	NM_000142	4D	204379_s_at
breast carcinoma amplified sequence 1	BCAS1	8537	NP_003648	NM_003657	4D	204378_at
Vprbinding protein	VprBP	9730	NP_055518	NM_014703	4D	204376_at
KHtype splicing regulatory protein (FUSE binding p	KHSRP	8570	NP_003676	NM_003685	40	2043/2_s_at
ATP/GTPbinding protein	HEAB	10978	NP_006822	NM_006831	4D	204370_at
general transcription factor IIIC, polypeptide 2,	GTF3C2	2976	NP_001512	NM_001521	4D	204366_s_at
TNF receptorassociated factor 5	TRAF5	7188	NP_004610	NM_004619	4D	204352_at
S100 calcium binding protein P	S100P	6286	NP_005971	NM_005980	4D	204351_at
collagen, type XVI, alpha 1	COL16A1	1307	NP_001847	NM_001856	4D	204345_at
zinc finger protein 202	ZNF202	7753	NP_003446	NM_003455	4D	204327_s_at
				NM_002450	4D	204326_x_at
natriuretic peptide receptor B/guanylate cyclase B	NPR2	4882	NP_003986	NM_003995	4D	204310_s_at
aldehyde dehydrogenase 6 family, member A1	ALDH6A1	4329	NP_005580	NM_005589	4D	204290_s_at
pim2 oncogene	PIM2	11040	NP_006866	NM_006875	4D	204269_at
S100 calcium binding protein A2	S100A2	6273	NP_005969	NM_005978	4D	204268_at
cyclindependent kinase 2	CDK2	1017	NP_001789	NM_001798	4D	204252_at
GULP, engulfment adaptor PTB domain containing 1	GULPI	51454	NP_057399	NM_016315	45	204235_S_at
chromosome 7 open reading frame 23	C7orf23	79161	NP_077291	NM_024315	#2	204215_at
solute carrier family 31 (copper transporters), me	SLC31A2	1318	NP_001851	NM_001860	4D	204204_at
BTB and CNC homology 1, basic leucine zipper trans	BACHI	571	NP_001177	NM_001186	40	204194_at
peptidylprolyl isomerase D (cyclophilin D)	PPID	5481	NP_005029	NM_005038	4D	204186_s_at

farnesyltransferase, CAAX box, beta	FNIB	2342	NP_002019	NM_002028	45	204/04_at
prostaglandinendoperoxide synthase 2 (prostaglandi	PIGS2	5/43	NF_000954	NIM_000903	5 5	204740_4
isoleucinetRNA synthetase	IARS	33/6	NP_038203	NIM 000072	ŧ	204749 st
(melanomaassoci		221	20202	LIVCIO WIN		20/7/// 6 24
chondroitin sulfate proteoglycan 4	CSPG4	1464	NP_001888	NM_001897	4D	204736_s_at
NCK adaptor protein 1	NCK1	4690	NP_006144	NM_006153	4D	204725_s_at
coagulation factor V (proaccelerin, labile factor)	F5	2153	NP_000121	NM_000130	4D	204713_s_at
hypothetical protein MGC29875	MGC29875	27042	NP_055203	NM_014388	4D	204700_x_at
cadherin 5, type 2, VEcadherin (vascular epitheliu	CDH5	1003	NP_001786	NM_001795	4D	204677_at
steroid5alphareductase, alpha polypeptide 1 (30xo5	SRD5A1	6715	NP_001038	NM_001047	4D	3 1
ring finger protein 24	RNF24	11237	NP_009150	NM_007219	4D	204668_at
adenosine deaminase	ADA	100	NP_000013	NM_000022	4D	204639_at
golgi SNAP receptor complex member 1	GOSR1	9527	NP_004862	NM_004871	4D	204630_s_at
nuclear receptor subfamily 4, group A, member 2	NR4A2	4929	NP_006177	NM_006186	4D	204622_x_at
adrenocortical dysplasia homolog (mouse)	ACD	65057	NP_075065	NM_022914	4D	204617_s_at
dickkopf homolog 1 (Xenopus laevis)	DKK1	22943	NP_036374	NM_012242	4D	204602_at
protein (peptidylprolyl cis/trans isomerase) NIMAi	PIN4	5303	NP_006214	NM_006223	4D	204571_x_at
KIAA0831	KIAA0831	22863	NP_055739	NM_014924	4D	204568_at
FK506 binding protein 5	FKBP5	2289	NP_004108	NM_004117	4D	204560_at
protein phosphatase 1, regulatory subunit 3D	PPPIR3D	5509	NP_006233	NM_006242	4D	204555_s_at
protein phosphatase 1, regulatory subunit 3D	PPP1R3D	5509	NP_006233	NM_006242	4D	204554_at
HermanskyPudlak syndrome 5	HPS5	11234	NP_009147	NM_007216	4D	204544_at
eukaryotic translation elongation factor 1 alpha 2	EEF1A2	1917	NP_001949	NM_001958	4D	204540_at
nuclear pore complex interacting protein	NPIP	9284	NP_008916	NM_006985	4D	204538_x_at
breast cancer 1, early onset	BRCA1	672	NP_009226	NM_007295	4D	204531_s_at
myosin VA (heavy polypeptide 12, myoxin)	MYO5A	4644	NP_000250	NM_000259	4D	204527_at
TBC1 domain family, member 8 (with GRAM domain)	TBC1D8	11138	NP_008994	NM_007063	40	204526_s_at
CDC7 cell division cycle 7 (S. cerevisiae)	CDC7	8317	NP_003494	NM_003503	40	204510_at
protein phosphatase 3 (formerly 2B), regulatory su	PPP3R1	5534	NP_000936	NM_000945	4D	204507_s_at
BH3 interacting domain death agonist	BID	637	NP_001187	NM_001196	4D	204493_at
transmembrane protein 15	TMEM15	22845	NP_055723	NM_014908	4D	204488_at
enolase 3 (beta, muscle)	ENO3	2027	NP_001967	NM_001976	4D	204483_at

clock homolog (mouse)	CLOCK	9575	NP_004889	NM_004898	#0	204980_at
neutrophil cytosolic factor 1 (47kDa, chronic gran	NCFI	4687		C97000 MN	3 3	204901 S at
myeloid cell nuclear differentiation antigen	MNDA	4332		NM_002432	4D	
pololike kinase 3 (Drosophila)	PLK3	1263	NP_004064	NM_004073	4D	204958_at
origin recognition complex, subunit 5like (yeast)	ORC5L	5001	NP_002544	NM_002553	4D	204957_at
intercellular adhesion molecule 3	ICAM3	3385	NP_002153	NM_002162	4D	204949_at
protein tyrosine phosphatase, receptor type, G	PTPRG	5793	NP_002832	NM_002841	4D	204944_at
inhibin, beta A (activin A, activin AB alpha polyp	INHBA	3624	NP_002183	NM_002192	4D	204926_at
tolllike receptor 2	TLR2	7097	NP_003255	NM_003264	4D	204924_at
myeloid/lymphoid or mixedlineage leukemia (trithor	MLLT3	4300	NP_004520	NM_004529	4D	204917_s_at
eukaryotic translation elongation factor 1 epsilon	EEF1E1	9521	NP_004271	NM_004280	4D	204905_s_at
amine oxidase, copper containing 3 (vascular adhes	AOC3	8639	NP_003725	NM_003734	4D	204894_s_at
HUS1 checkpoint homolog (S. pombe)	HUS1	3364	NP_004498	NM_004507	4D	204884 s_at
Rho GTPase activating protein 25	ARHGAP25	9938	NP_055697	NM_014882	4D	204882_at
immature colon carcinoma transcript 1	ICT1	3396	NP_001536	NM_001545	4D	204868_at
baculoviral IAP repeatcontaining 1	BIRCI	4671	NP_004527	NM_004536	4D	204861_s_at
baculoviral IAP repeatcontaining 1	BIRC1	4671	NP_004527	NM_004536	4D	204860_s_at
protein kinase, cAMPdependent, regulatory, type	PRKAR2A	5576	NP_004148	NM_004157	4D	204842_x_at
early endosome antigen 1, 162kD	EEA1	8411	NP_003557	NM_003566	4D	204840_s_at
processing of precursor 5, ribonuclease P/MRP subu	POP5	51367	NP_057002	NM_015918	4D	204839_at
mutL homolog 3 (E. coli)	MLH3	27030	NP_055196	NM_014381	4D	204838 s_at
major histocompatibility complex, class I, F	HLAF	3134	NP_061823	NM_018950	4D	204806_x_at
H1 histone family, member X	HIFX	8971	NP_006017	NM_006026	4D	204805_s_at
hypothetical protein FLJ13639	FLJ13639	79758	NP_078981	NM_024705	4D	204800_s_at
G proteincoupled receptor associated sorting prote	GPRASP1	9737	NP_055525	NM_014710	4D	204793_at
Fas (TNF receptor superfamily, member 6)	FAS	355	NP_000034	NM_000043	4D	204781 s_at
Fas (TNF receptor superfamily, member 6)	FAS	355	NP_000034	NM_000043	4D	204780_s_at
ecotropic viral integration site 2A	EVI2A	2123	NP_055025	NM_014210	4D	204774_at
transcription termination factor, RNA polymerase	TTF1	7270	NP_031370	NM_007344	4D	204 / /1_s_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT1	4521	NP_002443	NM_002452	4D	204766_s_at

vmaf musculoaponeurotic fibrosarcoma oncogene homo	MAFF	23764	NP_036455	NM_012323	4D	205193_at
retinitis pigmentosa 2 (Xlinked recessive)	RP2	6102	NP_008846	NM_006915	4D	205191_at
a disintegrin and metalloproteinase domain 8	ADAM8	101	NP_001100	NM_001109	4D	205179_s_at
CD58 antigen, (lymphocyte functionassociated antig	CD58	965	NP_0017/0	NM_0017/9	45	2051/3_x_at
leucine rich repeat neuronal 5	LRRN5	10446	NP_006329	NM_006338	4D	205154_at
chloride channel 4	CLCN4	1183	NP_001821	NM_001830	4D	205149_s_at
hypothetical protein FLJ22269	FLJ22269	84179	NP_115595	NM_032219	4D	205144_at
uronyl2sulfotransferase	UST	10090	NP_005706	NM_005715	4D	205139_s_at
heat shock 10kDa protein 1 (chaperonin 10)	HSPEI	3336	NP_002148	NM_002157	4D	205133_s_at
sarcoglycan, beta (43kDa dystrophinassociated glyc	SGCB	6443	NP_000223	NM_000232	4D	205121_at
formyl peptide receptor 1	FPR1	2357	NP_002020	NM_002029	4D	205119_s_at
syntaphilin	SNPH	9751	NP_055538	NM_014723	4D	205104_at
solute carrier family 26 (sulfate transporter), me	SLC26A2	1836	NP_000103	NM_000112	4D	205097_at
Nacetylglucosamine l phosphodiester alphaNacetylgluc	NAGPA	51172	NP_057340	NM_016256	4D	205090_s_at
interleukin 1, beta	ILIB	3553	NP_000567	NM_000576	4D	205067_at
AT rich interactive domain 4A (RBP1like)	ARID4A	5926	NP_002883	NM_002892	4D	205062_x_at
primase, polypeptide 1, 49kDa	PRIMI	5557	NP_000937	NM_000946	4D	205053_at
vkit HardyZuckerman 4 feline sarcoma viral oncogen	KIT	3815	NP_000213	NM_000222	4D	205051_s_at
				NM_003832	4D	205048_s_at
glucosamine (UDPNacetyl)2epimerase/Nacetylmannosam	GNE	10020	NP_005467	NM_005476	4D	205042_at
LSM6 homolog, U6 small nuclear RNA associated (S.	LSM6	11157	NP_009011	NM_007080	4 D	205036_at
cyclin E2	CCNE2	9134	NP_004693	NM_004702	4D	205034_at
transforming growth factor, alpha	TGFA	7039	NP_003227	NM_003236	4D	205016_at
transforming growth factor, alpha	TGFA	7039	NP_003227	NM_003236	4D	205015_s_at
trefoil factor 1 (breast cancer, estrogeninducible	TFF]	7031	NP_003216	NM_003225	4D	205009_at
Nmyristoyltransferase 2	NMT2	9397	NP_004799	NM_004808	4D	205006_s_at
Nmyristoyltransferase 2	NMT2	9397	NP_004799	NM_004808	4D	205005_s_at
cyclindependent kinase 5, regulatory subunit 1 (p3	CDK5R1	8851	NP_003876	NM_003885	4D	204996_s_at
integrin, beta 4	ITGB4	3691	NP_000204	NM_000213	4D	204989_s_at

butyrylcholinesterase	ВСНЕ	590	NP_000046	NM_000055	4D	205433_at
adaptorrelated protein complex 1, beta 1 subunit	APIBI	162	NP_001118	NM_001127	4D	205423_at
integrin, betalike 1 (with EGFlike repeat domains)	ITGBL1	9358	NP_004782	NM_004791	4D	205422_s_at
KIAA0672 gene product	KIAA0672	9912	NP_055674	NM_014859	4D	205414_s_at
sperm autoantigenic protein 17	SPA17	53340	NP_059121	NM_017425	4D	205406_s_at
interleukin 1 receptor, type II	IL1R2	7850	NP_004624	NM_004633	4D	205403_at
FXYD domain containing ion transport regulator (FXYDI	5348	NP_005022	NM_005031	4υ	205384_at
sarcolipin	SLN	6588	NP_003054	NM_003063	4D	205374_at
dihydrolipoamide branched chain transacylase E2	DBT	1629	NP_001909	NM_001918	4D	205371_s_at
glutamate receptor, ionotropic, AMPA 2	GRIA2	2891	NP_000817	NM_000826	4D	205358_at
thymosin, beta, identified in neuroblastoma cells	TMSNB	11013	NP_068832	NM_021992	4D	205347_s_at
ST3 betagalactoside alpha2,3sialyltransferase 2	ST3GAL2	6483	NP_008858	NM_006927	4D	205346_at
zinc finger and BTB domain containing 24	ZBTB24	9841	NP_055612	NM_014797	4D	205340_at
sorting nexin 4	SNX4	8723	NP_003785	NM_003794	4D	205329_s_at
activin A receptor, type II	ACVR2	92	NP_001607	NM_001616	4D	205327_s_at
kinesin family member 5A	KIF5A	3798	NP_004975	NM_004984	4D	205318_at
solute carrier family 15 (H+/peptide transporter),	SLC15A2	6565	NP_066568	NM_021082	4D	205317_s_at
				NM_012066	4D	205310_at
retinoblastomalike 1 (p107)	RBL1	5933	NP_899662	NM_183404	4D	205296_at
transcription factor AP2 gamma (activating enhance	TFAP2C	7022	NP_003213	NM_003222	4D	205286_at
FYN binding protein (FYB120/130)	FYB	2533	NP_001456	NM_001465	4D	205285_s_at
glycine receptor, beta	GLRB	2743	NP_000815	NM_000824	4D	205279_s_at
chromosome 21 open reading frame 5	C21orf5	9980	NP_005119	NM_005128	4D	205248_at
interleukin 1 receptor accessory protein	ILIRAP	3556	NP_002173	NM_002182	4D	205227_at
enoylCoenzyme A, hydratase/3hydroxyacyl Coenzyme A	ЕННАДН	1962	NP_001957	NM_001966	4D	205222_at
homogentisate 1,2dioxygenase (homogentisate oxidas	HGD	3081	NP_000178	NM_000187	4D	205221_at
polymerase (RNA) III (DNA directed) polypeptide F,	POLR3F	10621	NP_006457	NM_006466	4D	205218_at
apolipoprotein H (beta2glycoprotein I)	APOH	350	NP_000033	NM_000042	4D	205216_s_at
serine/threonine kinase 17b (apoptosisinducing)	STK17B	9262	NP_004217	NM_004226	4D	205214_at
vrel reticuloendotheliosis viral oncogene homolog	RELB	5971	NP_006500	NM_006509	4D	205205_at
phosphoserine phosphatase	PSPH	5723	NP_004568	NM_004577	4D	205194_at

betalipo						
proopiomelanocortin (adrenocorticotropin/	POMC	5443	NP_000930	NM_000939	4D	205720_at
bone marrow stromal cell antigen 1	BST1	683	NP_004325	NM_004334	4D	205715_at
interleukin 17 receptor	IL17R	23765	NP_055154	NM_014339	4D	205707_at
mitogenactivated protein kinase kinase 6	MAP2K6	5608	NP_002749	NM_002758	4D	205698_s_at
maternal G10 transcript	G10	8896	NP_003901	NM_003910	4D	205690 s_at
BCL2related protein A1	BCL2A1	597	NP_004040	NM_004049	4D	205681_at
major histocompatibility complex, class II, DO bet	HLADOB	3112	NP_002111	NM_002120	4D	205671_s_at
complement component 4 binding protein, alpha	C4BPA	722	NP_000706	NM_000715	4D	205654_at
small nuclear ribonucleoprotein polypeptide G	SNRPG	6637	NP_003087	NM_003096	4D	
centrosomal protein 1	CEP1	11064	NP_008949	NM_007018	4D	205642_at
cytidine deaminase	CDA	978	NP_001776	NM_001785	4D	
multimerin l	MMRNI	22915	NP_031377	NM_007351	4D	205612_at
homeo box B5	НОХВ5	3215	NP_002138	NM_002147	4D	205600_x_at
TRAF interacting protein	TRIP	10293	NP_005870	NM_005879	4D	205598_at
FGFR1 oncogene partner	FGFRIOP	111116	NP_008976	NM_007045	4D	205588_s_at
lysosomalassociated membrane protein 3	LAMP3	27074	NP_055213	NM_014398	4D	205569_at
carbohydrate (keratan sulfate Gal6) sulfotransfera	CHST1	8534	NP_003645	NM_003654	4D	205567_at
frataxin	FXN	2395	NP_000135	NM_000144	4D	205565_s_at
ribonuclease P/MRP 38kDa subunit	RPP38	10557	NP_006405	NM_006414	4D	205562_at
TNF receptorassociated factor 6	TRAF6	7189	NP_004611	NM_004620	4D	205558_at
deoxyribonuclease Ilike 3	DNASE1L3	1776	NP_004935	NM_004944	4D	205554_s_at
G1 to S phase transition 2	GSPT2	23708	NP_060564	NM_018094	4D	205541 s at
cadherin 6, type 2, Kcadherin (fetal kidney)	CDH6	1004	NP_004923	NM_004932	4D	205532_s_at
hyaluronan and proteoglycan link protein I	HAPLNI	1404	NP_001875	NM_001884	4D	205523_at
transcobalamin I (vitamin B12 binding protein, R	TCN1	6947	NP_001053	NM_001062	4D	205513_at
villin 1	VIL1	7429	NP_009058	NM_007127	4D	205506_at
KIAA0408	KIAA0408	9729	NP_055517	NM_014702	4D	205496_at
sorting nexin 15	SNX15	29907	NP_037438	NM_013306	4D	205482_x_at
adenosine A1 receptor	ADORAI	134	NP_000665	NM_000674	4D	205481_at
microfibrillarassociated protein 3 like	MFAP3L	9848	NP_067679	NM_021647	4D	205442_at
hypothetical protein FLJ22709	FLJ22709	79629	NP_078854	NM_024578	4D	205441_at
protein tyrosine phosphatase, nonreceptor type 21	PTPN21	11099	NP_008970	NM_007039	4D	205438_at
H2A histone family, member X	H2AFX	3014	NP_002096	NM_002105	4D	205436_s_at

				NM_014645	4D	206003_at
ELK4, ETSdomain protein (SRF accessory protein 1)	ELK4	2005	NP_001964	NM_001973	4D	205994_at
Tbox 2	TBX2	6909	NP_005985	NM_005994	4D	205993 s_at
CD1C antigen, c polypeptide	CDIC	911	NP_001756	NM_001765	4D	205987_at
klotho	KL	9365	NP_004786	NM_004795	4D	205978_at
histone 1, H4c	HIST1H4C	8364	NP_003533	NM_003542	4D	205967_at
basic leucine zipper transcription factor, ATFlike	BATF	10538	NP_006390	NM_006399	4D	205965_at
DnaJ (Hsp40) homolog, subfamily A, member 3	DNAJA3	9093	NP_005138	NM_005147	4D	205963_s_at
pyruvate dehydrogenase kinase, isoenzyme 4	PDK4	5166	NP_002603	NM_002612	4D	205960_at
plexin B3	PLXNB3	5365	NP_005384	NM_005393	4D	205957_at
carbonic anhydrase I	CAI	759	NP_001729	NM_001738	4D	205950_s_at
vanin 2	VNN2	8875	NP_004656	NM_004665	4D	205922_at
solute carrier family 6 (neurotransmitter transpor	SLC6A6	6533	NP_003034	NM_003043	4D	205921_s_at
solute carrier family 6 (neurotransmitter transpor	SLC6A6	6533	NP_003034	NM_003043	4D	205920_at
prepronociceptin	PNOC	5368	NP_006219	NM_006228	4D	205901_at
Jak and microtubule interacting protein 2	KIAA0555	9832	NP_055605	NM_014790	4D	205888_s_at
leukemia inhibitory factor receptor	LIFR	3977	NP_002301	NM_002310	4D	205876_at
AT rich interactive domain 3A (BRIGHT like)	ARID3A	1820	NP_005215	NM_005224	4D	205865_at
solute carrier family 7 (cationic amino acid trans	SLC7A4	6545	NP_004164	NM_004173	4D	205864_at
solute carrier family 18 (vesicular monoamine), me	SLC18A2	6571	NP_003045	NM_003054	4D	205857_at
polymerase (DNA directed), gamma 2, accessory subu	POLG2	11232	NP_009146	NM_007215	4D	205811_at
tuftelin 1	TUFT1	7286	NP_064512	NM_020127	4D	205807_s_at
TRAF3 interacting Jun Nterminal kinase (JNK) activat	T3JAM	80342	NP_079504	NM_025228	4D	205804_s_at
tyrosine kinase, nonreceptor, 1	TNKI	8711	NP_003976	NM_003985	4D	205793_x_at
receptor (calcitonin) activity modifying protein 2	RAMP2	10266	NP_005845	NM_005854	4D	205779_at
coagulation factor XII (Hageman factor)	FI2	2161	NP_000496	NM_000505	4D	205774_at
DEAD (AspGluAlaAsp) box polypeptide 18	DDX18	8886	NP_006764	NM_006773	4D	205763_s_at
coagulation factor VIII, procoagulant component (h	F8	2157	NP_000123	NM_000132	4D	205756_s_at
biphenyl hydrolaselike (serine hydrolase; breast e	BPHL	670	NP_004323	NM_004332	4D	205750_at
oncostatin M receptor	OSMR	9180	NP_003990	NM_003999	4D	205729_at
ciliary neurotrophic factor receptor	CNTFR	1271	NP_001833	NM_001842	4D	205723_at

miltogenactivated protein kinase kinase kinase kina	MAP471	1110+	141 007112	17111 00/101		and the second s
wante and the state of the stat	MADAVI	11107	C11000 dlv	NM 007181	45	206296 x at
teratocarcinomaderived growth factor 1	TDGFI	6997	NP 003203	NM_003212	4D	206286_s_at
arginine vasopressin receptor 1A	AVPRIA	552	NP_000697	NM_000706	4D	206250_x_at
YY1 associated factor 2	YAF2	10138	NP_005739	NM_005748	4D	206238_s_at
G proteincoupled receptor 4	GPR4	2828	NP_005273	NM_005282	4D	a
paired box gene 2	PAX2	5076	NP_003979	NM_003988	4D	
tumor necrosis factor receptor superfamily, member	TNFRSF10C	8794		NM_003841	45	at
carbonic anhydrase IV	CA4	762	NP_000708	NM_000717	100	206209 s_at
carbonic anhydrase IV	CA4	762	NP_000708	NM_000717	4D	206208_at
Mphase phosphoprotein 9	MPHOSPH9	10198	NP_073619	NM_022782	4D	206205_at
unc5 homolog C (C. elegans)	UNC5C	8633	NP_003719	NM_003728	4D	206189_at
crystallin, beta B1	CRYBB1	1414	NP_001878	NM_001887	4D	206185_at
arginase, liver	ARG1	383	NP_000036	NM_000045	4D	206177 s_at
adrenergic, beta2, receptor, surface	ADRB2	154	NP_000015	NM_000024	4D	206170_at
pentaxinrelated gene, rapidly induced by IL1 beta	PTX3	5806	NP_002843	NM_002852	4D	206157_at
LIM homeobox 2	LHX2	9355	NP_004780	NM_004789	4D	206140_at
frizzled homolog 5 (Drosophila)	FZD5	7855	NP_003459	NM_003468	4D	206136_at
arylsulfatase B	ARSB	411	NP_000037	NM_000046	4D	206129 s_at
early growth response 3	EGR3	1960	NP_004421	NM_004430	4D	206115_at
EPH receptor A4	EPHA4	2043	NP_004429	NM_004438	4D	206114_at
histone 1, H3h	HIST1H3H	8357	NP_003527	NM_003536	40	206110_at
	RAC3	5881	NP_005043	NM_005052	4D	206103_at
solute carrier family 22 (organic cation transport	SLC22A1LS	5003	NP_009036	NM_007105	4D	206097_at
NELlike I (chicken)	NELL1	4745	NP_006148	NM_006157	4D	206089_at
high mobility group AThook	HMGA1	3159	NP_002122	NM_002131	4D	206074_s_at
urocortin	UCN	7349	NP_003344	NM_003353	4D	206072_at
RAD51 homolog C (S. cerevisiae)	RAD51C	5889	NP_002867	NM_002876	4D	206066_s_at
small nuclear ribonucleoprotein polypeptide A'	SNRPAI	6627	NP_003081	NM_003090	4D	206055_s_at
zinc finger protein 510	ZNF510	22869	NP_055745	NM_014930	4D	206053_at
selectin P (granule membrane protein 140kDa, antig	SELP	6403	NP_002996	NM_003005	4D	
nucleolar protein 4	NOL4	8715	NP_003778	NM_003787	4D	206045_s_at
RAB33A, member RAS oncogene family	RAB33A	9363	NP_004785	NM_004794	4D	206039_at
aspartoacylase (aminoacylase 2, Canavan disease)	ASPA	443	NP_000040	NM_000049	4D	206030_at

mitogenactivated protein kinase kinase kinase kina	MAP4K4	9448	NP_004825	NM_004834	140	2003/1_S_at
	NAADAKA	0440	NID ONASOS	NIM 00/1827	3	ا :
SMA3	SMA3	10571	NP 006771	NM 006780	4D	206565 x at
nucleoporin 155kDa	NUP155	9631	NP_004289	NM_004298	4D	206550_s_at
SWI/SNF related, matrix associated, actin dependen	SMARCBI	6598	NP_003064	NM_0030/3	4D	206032_at
4aminobutyrate aminotransferase	ABAT	18	NP_000654	NM_000663	4D	206527_at
maltaseglucoamylase (alphaglucosidase)	MGAM	8972	NP_004659	NM_004668	4D	206522_at
sine oculis homeobox homolog 2 (Drosophila)	SIX2	10736	NP_058628	NM_016932	4D	206511_s_at
oculocutaneous albinism II (pinkeye dilution homol	OCA2	4948	NP_000266	NM_000275	4D	206498_at
MBD2 (methylCpGbinding protein)interacting zinc fi	MIZF	25988	NP_056332	NM_015517	4D	206495_s_at
LIM domain binding 2	LDB2	9079	NP_001281	NM_001290	4D	206481 s_at
transducinlike enhancer of split 3 (E(spl) homolog	TLE3	7090	NP_005069	NM_005078	4D	206472 s_at
neurotrophic tyrosine kinase, receptor, type 3	NTRK3	4916	NP_002521	NM_002530	†Đ	206462_s_at
rhodopsin (opsin 2, rod pigment) (retinitis pigmen	RHO	6010	NP_000530	NM_000539	4D	206454_s_at
lin7 homolog A (C. elegans)	LIN7A	8825	NP_004655	NM_004664	4D	206440_at
coagulation factor II (thrombin) receptorlike 1	F2RL1	2150	NP_005233	NM_005242	4D	206429_at
melanA	MLANA	2315	NP_005502	NM_005511	4D	206427_s_at
cyclic nucleotide gated channel alpha 1	CNGAI	1259	NP_000078	NM_000087	4D	206417_at
development and differentiation enhancing factor 2	DDEF2	8853	NP_003878	NM_003887	4D	206414_s_at
ubiquitin specific protease 6 (Tre2 oncogene)	USP6	9098	NP_004496	NM_004505	4D	206405_x_at
CD19 antigen	CD19	930	NP_001761	NM_001770	4D	206398_s_at
retinoic acid receptor responder (tazarotene induc	RARRESI	5918	NP_002879	NM_002888	4D	206392_s_at
phosphodiesterase 3A, cGMPinhibited	PDE3A	5139	NP_000912	NM_000921	4D	206389_s_at
serine (or cysteine) proteinase inhibitor, clade A	SERPINA7	6906	NP_000345	NM_000354	4D	206386_at
brainderived neurotrophic factor	BDNF	627	NP_001700	NM_001709	4D	206382_s_at
myogenic factor 6 (herculin)	MYF6	4618	NP_002460	NM_002469	4D	206372_at
chemokine (C motif) ligand 1	XCL1	6375	NP_002986	NM_002995	4D	206365_at
suppressor of cytokine signaling 3	SOCS3	9021	NP_003946	NM_003955	4D	206359_at
amyloid P component, serum	APCS	325	NP_001630	NM_001639	4D	206350_at
iduronate 2sulfatase (Hunter syndrome)	IDS	3423	NP_006114	NM_006123	4D	206342_x_at
chemokine (CXC motif) ligand 6 (granulocyte chemot	CXCL6	6372	NP_002984	NM_002993	4D	206336_at
ryanodine receptor 3	RYR3	6263	NP_001027	NM_001036	4D	206306_at

142 NM, 004387 NP, 004378 1482 NKX2rs NKX2rsmscription factor related, locus 5 (Drosoph 4D) NM, 004361 NP 066246 S5634 FL20344 Sypothetical protein Fl20344 Sypoth							
NM_004387 NP_004378 I482 NKX25	zinc finger protein 430	ZNF430	80264		NM_025189	4D	206829_x_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_066179 23643 LY96 NM_015364 NP_004552 5017 OVOL1 NM_004561 NP_003846 8809 IL18R1 NM_004561 NP_005579 9719 ADAMTSL2 NM_004900 NP_005599 9719 ADAMTSL2 NM_004900 NP_005599 9719 ADAMTSL2 NM_004900 NP_005599 9719 ADAMTSL2 NM_006667 NP_006898 3357 HTR2B NM_0020521 NP_006392 57136 C20orf3 NM_006693 NP_006392 57136 C20orf3 NM_006684 NP_006399 57136 CPSF4 NM_00688 NP_006392 57136 C20orf3 NM_005891 NP_006392 57136 CPSF4 NM_005864 NP_003982 1910 EDNRB NM_018953 NP_005826 8690 JRKL	peripheral myelin protein 2	PMP2	5375		NM_002677	4D	206826_at
NM_004387 NP_004378 I482 NKX25 NM_01776 NP_060246 55634 FLJ20344 NM_015364 NP_066179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_004900 NP_003846 8809 IL18R1 NM_004900 NP_004891 9582 APOBEC3B NM_004900 NP_004891 9582 APOBEC3B NM_004900 NP_004891 9582 APOBEC3B NM_004900 NP_004891 9582 APOBEC3B NM_006384 NP_006392 57136 C20orf3 NM_006693 NP_066392 57136 C20orf3 NM_006684 NP_006379 10524 HTATIP NM_006388 NP_003982 1910 EDNRB NM_004201 NP_003982 1910 EDNRB NM_004202 NP_0078995 79774 GRTPI NM_004202 NP_001603 36 ACRVI	sperm associated antigen 8	SPAG8	26206		NM_012436	4D	206816_s_at
NM_004387 NP_004378 I482 NKX25 NM_01776 NP_060246 55634 FLJ20344 NM_01776 NP_060246 55634 FLJ20344 NM_015364 NP_066179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_004561 NP_003846 8809 IL18R1 NM_004900 NP_003846 8809 IL18R1 NM_004900 NP_003846 9719 ADAMTSL2 NM_004900 NP_00385 3357 HTR2B NM_004900 NP_00689 3357 HTR2B NM_0016384 NP_006392 57136 C20orf3 NM_006693 NP_066392 57136 C20orf3 NM_006388 NP_006379 10524 HTATIP NM_006389 NP_003982 1910 EDNRB NM_012307 NP_036439 23136 EPB41L3 NM_01864 NP_003982 1137 CHRNA4 NM_004202 NP_001893 3741 KCNA5					NM_005758	4D	206809_s_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_060246 55634 FLJ20344 NM_015364 NP_060252 5017 OVOL1 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_004900 NP_0055509 9719 ADAMTSL2 NM_004900 NP_0055509 9719 ADAMTSL2 NM_004900 NP_0055509 9719 ADAMTSL2 NM_004900 NP_0055509 9719 ADAMTSL2 NM_00467 NP_005392 57136 C20orf3 NM_005381 NP_006392 57136 C20orf3 NM_0050693 NP_006379 10524 HTATIP NM_005384 NP_003982 1910 EDNRB NM_015864 NP_003962 9750 C6orf32 NM_003971 NP_003962 32136 EPB41L3 NM_005234 NP_0061826 3222 HOXC5	prodynorphin	PDYN	5173	NP_077722	NM_024411	4D	206803_at
NM_004387 NP_004378 I482 NKX25 NM_01776 NP_060246 55634 FLJ20344 NM_015364 NP_060246 55634 FLJ20344 NM_015364 NP_060252 5017 OVOL1 NM_004561 NP_004552 5017 OVOL1 NM_004900 NP_003846 8809 IL18R1 NM_016694 NP_055509 9719 ADAMTSI.2 NM_004900 NP_0055509 9719 ADAMTSI.2 NM_016384 NP_006858 3357 HTR2B NM_016384 NP_006392 57136 C20orf3 NM_006693 NP_006379 10524 HTATIP NM_003864 NP_003982 1910 EDNRB NM_015864 NP_0036439 23136 EPB41L3 NM_015864 NP_003763 8690 JRKL NM_003971 NP_003962 3913 CHRNA4 NM_0040202 NP_004193 9087 TMSB4Y NM_001612 NP_001603 56 ACRV1	WNT1 inducible signaling pathway protein 1	WISPI	8840	NP_003873	NM_003882	4D	206796_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_004900 NP_004891 9582 APOBEC3B NM_004900 NP_004891 9582 APOBEC3B NM_004900 NP_006858 3357 HTR2B NM_016384 NP_006392 57136 C20orf3 NM_016384 NP_006392 57136 C20orf3 NM_016384 NP_006379 10524 HTATIP NM_006693 NP_006379 10524 HTATIP NM_013864 NP_003982 1910 EDNRB NM_012307 NP_036439 23136 EPB41L3 NM_018953 NP_03763 8690 JRKL NM_018953 NP_0361826 3222 HOXC5 NM_003971 NP_003962 9043 SPAG9	verba erythroblastic leukemia viral oncogene homol	ERBB4	2066	NP_005226	NM_005235	40	1
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_065179 23643 LY96 NM_004561 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_000867 NP_000858 3357 HTR2B NM_016384 NP_00684 10898 CPSF4 NM_0025024 NP_00684 10898 CPSF4 NM_006693 NP_00684 10898 CPSF4 NM_0015864 NP_003982 1910 EDNRB NM_012307 NP_0036439 23136 EPB41L3 NM_018953 NP_007895 79774 GRTP1 NM_018953 NP_0061826 3222 HOXC5 NM_004202 NP_003962 9043 SPAG9	killer cell lectinlike receptor subfamily C, membe	KLRC2	3822	NP_002251	NM_002260	40	206/85_s_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_065179 23643 FLJ20344 NM_001856 NP_004891 23643 Ly96 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_0055509 9719 ADAMTSL2 NM_0016384 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_025024 NP_065392 57136 C20orf3 NM_006693 NP_066379 10524 HTATIP NM_003991 NP_063982 1910 EDNRB NM_015864 NP_036439 23136 EPB41L3 NM_008772 NP_036439 23136 EPB41L3 NM_018953 NP_003763 8690 JRKL NM_002234 NP_003962 9043 SPAG9	acrosomal vesicle protein 1	ACRV1	56	NP_001603	NM_001612	4D	206776_x_at
NM_004387 NP_004378 I482 NKX25 0 NM_017776 NP_060246 55634 FLJ20344 0 NM_015364 NP_056179 23643 LY96 0 NM_004561 NP_004552 5017 OVOL1 0 NM_004561 NP_003846 8809 IL18R1 0 NM_014694 NP_055509 9719 ADAMTSL2 0 NM_004900 NP_004891 9582 APOBEC3B 0 NM_004900 NP_000858 3357 HTR2B 0 NM_006693 NP_0065392 57136 C20orf3 0 NM_006693 NP_006379 10524 HTATIP 0 NM_00688 NP_003982 1910 EDNRB 0 NM_013864 NP_003982 1910 EDNRB 0 NM_012307 NP_036439 23136 EPB41L3 0 NM_013953 NP_003763 8690 JRKL 0 NM_000774 NP_0061826 3222 HOXC5	thymosin, beta 4, Ylinked	TMSB4Y	9087	NP_004193	NM_004202	4D	at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_017364 NP_065179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_003867 NP_004891 9582 APOBEC3B NM_016384 NP_0065392 57136 C20orf3 NM_020531 NP_065392 57136 C20orf3 NM_006693 NP_00684 10898 CPSF4 NM_003991 NP_003982 1910 EDNRB NM_012864 NP_036439 23136 EPB41L3 NM_018953 NP_003763 8690 JRKL NM_003971 NP_003962 9043 SPAG9	potassium voltagegated channel, shakerrelated subf	KCNA5	3741	NP_002225	NM_002234	40	206/62_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_017776 NP_060246 55634 FLJ20344 NM_013364 NP_066179 23643 LY96 NM_004561 NP_003845 8809 IL18R1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_025024 NP_066392 57136 C20orf3 NM_006693 NP_006379 10898 CPSF4 NM_003991 NP_003982 1910 EDNRB NM_012307 NP_036439 23136 EPB41L3 NM_018953 NP_003763 8690 JRKL NM_000744 NP_0061826 3222 HOXC5	sperm associated antigen 9	SPAG9	9043	NP_003962	NM_003971	4D	206748_s_at
NM_004387 NP_004378 I 482 NKX25 NM_01776 NP_060246 55634 FLJ20344 NM_015364 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_0065392 57136 C20orf3 NM_025024 NP_065392 57136 C20orf3 NM_006693 NP_006379 10524 HTATIP NM_003991 NP_003982 1910 EDNRB NM_015864 NP_036439 23136 EPB41L3 NM_003772 NP_036439 23136 EPB41L3 NM_000744 NP_000735 1137 CHRNA4	homeo box C5	HOXC5	3222	NP_061826	NM_018953	4D	206739_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_011776 NP_060246 55634 FLJ20344 NM_0115364 NP_065179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_006693 NP_065392 57136 C20orf3 NM_006693 NP_00684 10898 CPSF4 NM_015864 NP_003982 1910 EDNRB NM_015864 NP_036439 23136 EPB41L3 NM_024719 NP_036439 23136 EPB41L3 NM_03772 NP_03763 8690 JRKL	cholinergic receptor, nicotinic, alpha polypeptide	CHRNA4	1137	NP_000735	NM_000744	4D	206735_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_066179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_000858 3357 HTR2B NM_025024 NP_065392 57136 C20orf3 NM_006693 NP_066399 10898 CPSF4 NM_003991 NP_00684 10898 CPSF4 NM_013864 NP_036439 23136 EDNRB NM_012307 NP_036439 23136 EPB41L3 NM_024719 NP_078995 79774 GRTP1	jerky homologlike (mouse)	JRKL	8690	NP_003763	NM_003772	4D	206734_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_066179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_0055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_000858 3357 HTR2B NM_025024 NP_065392 57136 C20orf3 NM_006693 NP_00684 10898 CPSF4 NM_003991 NP_006379 10524 HTATIP NM_015864 NP_056948 9750 C6orf32 NM_012307 NP_036439 23136 EPB41L3	- 1	GRTPI	79774	NP_078995	NM_024719	4D	206712_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_014694 NP_003846 8809 IL18R1 NM_004900 NP_004891 9582 APOBEC3B NM_006384 NP_000858 3357 HTR2B NM_025024 NP_065392 57136 C20orf3 NM_006388 NP_006379 10524 HTATIP NM_015864 NP_056948 9750 C6orf32		EPB41L3	23136	NP_036439	NM_012307	4D	206710 s_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_004900 NP_004891 9582 APOBEC3B NM_000867 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_006693 NP_006379 10524 HTATIP NM_003991 NP_003982 1910 EDNRB	chromosome 6 open reading frame 32	C6orf32	9750	NP_056948	NM_015864	4D	206707_x_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_00667 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_006693 NP_006379 10524 HTATIP	endothelin receptor type B	EDNRB	1910	NP_003982	NM_003991	4D	206701_x_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_000867 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_025024 NP_006693 NP_006684 10898 CPSF4	HIV1 Tat interacting protein, 60kDa	HTATIP	10524		NM_006388	4D	206689_x_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_00667 NP_000858 3357 HTR2B NM_016384 NP_065392 57136 C20orf3 NM_025024 OVOL1 C20orf3	cleavage and polyadenylation specific factor 4, 30	CPSF4	10898	NP_006684	NM_006693	4D	206688_s_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_016384 NP_00858 3357 HTR2B NM_020531 NP_065392 57136 C20orf3	- (NM_025024	4D	206659_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_000867 NP_000858 3357 HTR2B NM_016384 HTR2B HTR2B	chromosome 20 open reading frame 3	C20orf3	57136	NP_065392	NM_020531	4D	206656_s_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B NM_000867 NP_000858 3357 HTR2B					NM_016384	4D	206652_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2 NM_004900 NP_004891 9582 APOBEC3B	5hydroxytryptamine (serotonin) receptor 2B	HTR2B	3357	NP_000858	NM_000867	4D	206638_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1 NM_014694 NP_055509 9719 ADAMTSL2	apolipoprotein B mRNA editing enzyme, catalytic po	APOBEC3B	9582	NP_004891	NM_004900	4D	206632_s_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1 NM_003855 NP_003846 8809 IL18R1	ADAMTSlike 2	ADAMTSL2	9719	NP_055509	NM_014694	4D	206629_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96 NM_004561 NP_004552 5017 OVOL1	interleukin 18 receptor 1	IL18R1	8809	NP_003846	NM_003855	4D	206618_at
NM_004387 NP_004378 1482 NKX25 NM_017776 NP_060246 55634 FLJ20344 NM_015364 NP_056179 23643 LY96	ovolike I(Drosophila)	OVOLI	5017		NM_004561	4D	206604_at
NM_004387 NP_004378 I482 NKX25 NM_017776 NP_060246 55634 FLJ20344	lymphocyte antigen 96	LY96	23643		NM_015364	4D	206584_at
NM_004387 NP_004378 1482 NKX25	hypothetical protein FLJ20344	FLJ20344	55634	NP_060246	NM_017776	4D	206583_at
	NK2 transcription factor related. locus 5 (Drosonh	NKX25	1482	NP_004378	NM_004387	4D	206578_at

206830_at	4D	NM_022058	NP_071341	57282	SLC4A10	solute carrier family 4, sodium bicarbonate transp
206832_s_at	4D	NM_004186	NP_004177	6405	SEMA3F	sema domain, immunoglobulin domain (Ig), short bas
206840_at	4D	NM_001133	NP_001124	173	AFM	afamin
206841_at	4D	NM_006205	961900_4N	5149	PDE6H	phosphodiesterase 6H, cGMPspecific, cone, gamma
206844_at	4D	NM_003837	NP_003828	8789	FBP2	fructose1,6bisphosphatase 2
206848_at	4D	NM_006896	NP_008827	3204	НОХА7	homeo box A7
206860_s_at	4D	NM_019005	NP_061878	54468	FLJ20323	hypothetical protein FLJ20323
206864_s_at	4D	NM_003806	NP_003797	8739	HRK	harakiri, BCL2 interacting protein (contains only
206866_at	4D	NM_001794	NP_001785	1002	CDH4	cadherin 4, type 1, Rcadherin (retinal)
206875_s_at	4D	NM_014720	NP_055535	9748	SLK	STE20like kinase (yeast)
206877_at	4D	NM_002357	NP_002348	4084	MXDI	MAX dimerization protein 1
206893_at	4D	NM_002968	NP_002959	6299	SALLI	sallike I (Drosophila)
206911_at	4D	NM_005082	NP_005073	7706	TRIM25	tripartite motifcontaining 25
206914_at	4D	NM_019604	NP_062550	56253	CRTAM	classI MHCrestricted T cell associated molecule
206920_s_at	4D	NM_001499	NP_001490	2733	GLEIL	GLE1 RNA export mediatorlike (yeast)
206923_at	4D	NM_002737	NP_002728	5578	PRKCA	protein kinase C, alpha
206941_x_at	4D	NM_012431	NP_036563	9723	SEMA3E	sema domain, immunoglobulin domain (lg), short bas
206942_s_at	4D	NM_002674	NP_002665	5367	РМСН	promelaninconcentrating hormone
206949_s_at	4D	NM_014328	NP_055143	23623	RUSC1	RUN and SH3 domain containing 1
206952_at	4D	NM_000151	NP_000142	2538	G6PC	glucose6phosphatase, catalytic (glycogen storage d
206970_at	4D	NM_005076	NP_005067	6900	CNTN2	contactin 2 (axonal)
206973_at	4D	NM_003625	NP_003616	8499	PPFIA2	protein tyrosine phosphatase, receptor type, f pol
206983_at	4D	NM_004367	NP_004358	1235	CCR6	chemokine (CC motif) receptor 6
206995_x_at	4D	NM_003693	NP_003684	8578	SCARFI	scavenger receptor class F, member 1
207001_x_at	4D	NM_004089	NP_004080	1831	DSIPI	delta sleep inducing peptide, immunoreactor
207006_s_at	4D	NM_013301	NP_037433	29903	HSU79303	protein predicted by clone 23882
207010_at	4D	NM_000812	NP_000803	2560	GABRB1	gammaaminobutyric acid (GABA) A receptor, beta 1
207018_s_at	4D	NM_004163	NP_004154	5874	RAB27B	RAB27B, member RAS oncogene family
207021_at	4D	NM_007009	NP_008940	11055	ZPBP	zona pellucida binding protein
207028_at	4D	NM_006316				
207036_x_at	4D	NM_000836	NP_000827	2906	GRIN2D	glutamate receptor, ionotropic, Nmethyl

carbonic anhydrase VA, mitochondrial	CA5A	/63	NP_001/30	MNI_001/39	40	20/421_dt
proprotein convertase subtilisin/kexin type 6	PCSK6	5046	NP_002561	NM_0025/0	B E	207414_S_at
homeo box D13	HOXD13	3239	NP 000514	NM_000523	4D	
glycoprotein lb (platelet), alpha polypeptide	GPIBA	2811	NP_000164	NM_000173	4D	207389_at
rhomboid, veinletlike I (Drosophila)	RHBDL1	9028	NP_003952	NM_003961	4D	207383_s_at
chromosome X open reading frame 2	CXorf2	1527	NP_001577	NM_001586	4D	207364_at
transforming growth factor, beta receptor II (70/8	TGFBR2	7048	NP_003233	NM_003242	4D	207334_s_at
melanoma antigen family A, I (directs expression o	MAGEAI	4100	NP_004979	NM_004988	#	20/222_x_dt
5hydroxytryptamine (serotonin) receptor 2C	HTR2C	3358	NP_000859	NM_000868	1 1 1 1	207307_at
matrix metalloproteinaselike 1	MMPL1	4328	NP_004133	NM_004142	4D	207289_at
KIAA0635	KIAA0635	9662	NP_079285	NM_025009	4D	207286_at
				NM_016365	4D	207279_s_at
solute carrier family 15 (oligopeptide transporter	SLC15A1	6564	NP_005064	NM_005073	4D	207254_at
ubinuclein l	UBNI	29855	NP_058632	NM_016936	4D	207253_s_at
meprin A, beta	MEPIB	4225	NP_005916	NM_005925	4D	207251_at
zinc finger DAZ interacting protein 3	DZIP3	9666	NP_055463	NM_014648	4D	207231_at
arylalkylamine Nacetyltransferase	AANAT	15	NP_001079	NM_001088	4D	207225_at
arachidonate 12lipoxygenase	ALOX12	239	NP_000688	NM_000697	4D	207206_s_at
carcinoembryonic antigenrelated cell adhesion mole	CEACAM4	1009	1001000	11111_OOLO17		***************************************
iciai Aizheilher antigen	CDACANA	1000	808100 div	NM 001817	40	207205 at
fetal Alabaimon articles	FAI 7	2186	NP 004450	NM 004459	4D	
immunoalohulin supostomilis monto.	IGSF2	8656	NP 004249	NM 004258	4D	207167 at
apolipoprotein B mRNA editing enzyme, catalytic	APOBECI	339	NP_001635	NM_001644	4D	207158_at
neurotrophic tyrosine kinase, receptor, type 2	NTRK2	4915	NP_006171	NM_006180	4D	207152_at
sulfotransferase family, cytosolic, 1A, phenolpref	SULT1A2	6799	NP_001045	NM_001054	4D	207122_x_at
solute carrier family 17 (sodium phosphate), membe	SLC17A2	10246	NP_005826	NM_005835	4D	20/09/_s_at
serum amyloid A4, constitutive	SAA4	6291	NP_006503	NM_006512	4D	207096_at
interleukin 18 receptor accessory protein	IL18RAP	8807	NP_003844	NM_003853	4D	207072_at
amine oxidase, copper containing 2 (retinaspecific	AOC2	314	NP_033720	NM_009590	4D	207064_s_at
Gprotein coupled receptor 37 like 1	GPR37L1	9283	NP_004758	NM_004767	4D	207055_at
mannanbinding lectin serine protease 2	MASP2	10747	NP_006601	NM_006610	4D	207041_at
Daspartate						
						-

keratin, hair, basic, 3	KRTHB3	3889	NP_002273	NM_002282	40	20/669_at
mitogenactivated protein kinase kinase 3	MAP2K3	5606	NP_002747	NM_002756	40	11
a disintegrin and metalloproteinase domain 21		8747	NP 003804	NM_003813	4D	at
Tbox 1	TBXI	6899	NP_005983	NM_005992	4D	
SH3 multiple domains 1	SH3MD1	9644	NP_055446	NM_014631	40	20/661_s_at
myelinassociated oligodendrocyte basic protein	MOBP	4336		NM_006501	4D	207659_s_at
transportin l	TNPO1	3842	NP_002261	NM_002270	4D	207657_x_at
Bcell linker	BLNK	29760	NP_037446	NM_013314	4D	207655_s_at
keratin, hair, acidic, 7	KRTHA7	8688	NP_003761	NM_003770	40	207649 at
tumor necrosis factor receptor superfamily, member	TNFRSF1A	7132	NP_001056	NM_001065	4D	1 1
protease, serine, 7 (enterokinase)	PRSS7	5651	NP_002763	NM_002772	4D	207638_at
serine (or cysteine) proteinase inhibitor, clade I	SERPINI2	5276	NP_006208	NM_006217	4D	207636_at
corebinding factor, runt domain, alpha subunit 2;	CBFA2T2	9139	NP_005084	NM_005093	4D	207625_s_at
hypocretin (orexin) receptor 1	HCRTR1	3061	NP_001516	NM_001525	40	20/619_at
calcium/calmodulindependent protein kinase (CaM ki	CAMK2A	815	NP_057065	NM_015981	ŧ	
sulfotransferase family, cytosolic, 1B, member 1	SULTIBI	27284	NP_055280	NM_014465	4D	207601_at
chromosome 1 open reading frame 38	Clorf38	9473	NP_004839	NM_004848	4D	207571_x_at
zinc finger protein 261	ZNF261	9203	NP_005087	NM_005096	4D	20/559_s_at
trophoblastlympho						
membrane cofactor protein (CD46.	MCP	4179	NP_002380	NM_002389	4D	207549_x_at
numb homolog (Drosophila)	NUMB	8650	NP_003735	NM_003744	4D	207545_s_at
exosome component 10	EXOSC10	5394	NP_002676	NM_002685	4D	207541 s_at
defensin, alpha 5, Paneth cellspecific	DEFA5	1670	NP_066290	NM_021010	4D	207529_at
bradykinin receptor B1	BDKRB1	623	NP_000701	NM_000710	4D	207510_at
TBPinteracting protein	0A	55832	NP_060918	NM_018448	4D	207483_s_at
motilin	MLN	4295	NP_002409	NM_002418	4D	207473_at
hypothetical protein DKFZp566H0824	DKFZp566H082 4	54744	NP_060005	NM_01/335	5	20/4/0_at
galanin	GAL	51083	NP_057057	NM_015973	8 8	20/466_at
glycophorin B (includes Ss blood group)	GYPB	2994	NP_002091	NM_002100	4D	207459_x_at
chemokine (CC motif) receptor 9	CCR9	10803	NP_006632	NM_006641	4D	207445 s_at
solute carrier family 35 (UDPgalactose transporter	SLC35A2	7355	NP_005651	NM_005660	4D	207439 s_at
solute carrier family 22 (organic cation transport	SLC22A2	6582	NP_003049	NM_003058	4D	207429_at

paired box gene 8	PAX8	7849	NP_054698	NM_013992	4D	20/924_x_at
cytochrome P450, family 2, subfamily F, polypeptid		1572	NP_000765	NM_000774	4D	207913_at
				NM_004940	4D	207888_at
nuclear VCPlike	NVL	4931	NP_002524	NM_002533	4D	207877_s_at
seizure related 6 homolog (mouse)like	SEZ6L	23544	NP_066938	NM_021115	4D	207873_x_at
leukocyte immunoglobulinlike receptor, subfamily A	LILRA2	111027	NP_006857	NM_006866	4D	207857_at
chemokine (CXC motif) ligand 3	CXCL3	2921	NP_002081	NM_002090	4D	207850_at
interleukin 13	IL13	3596	NP_002179	NM_002188	4D	207844_at
cytochrome b5-	CYB5	1528	NP_001905	NM_001914	4D	207843_x_at
fibulin 1	FBLN1	2192	NP_006478	NM_006487	4D	207835_at
BCL2/adenovirus E1B 19kDa interacting protein 1	BNIP1	662	NP_053581	NM_013978	4D	207829_s_at
inhibitor of DNA binding 3, dominant negative heli	ID3	3399	NP_002158	NM_002167	4D	207826_s_at
defensin, alpha 6, Paneth cellspecific	DEFA6	1671	NP_001917	NM_001926	4D	207814_at
ferredoxin reductase	FDXR	2232	NP_004101	NM_004110	4D	207813 s_at
keratin 12 (Meesmann corneal dystrophy)	KRT12	3859	NP_000214	NM_000223	4D	207811_at
ataxin 2like	ATXN2L	111273	NP_059867	NM_017492	4D	207798_s_at
RABIA, member RAS oncogene family	RAB1A	5861	NP_004152	NM_004161	4D	207791_s_at
keratin, hair, acidic, 3B	KRTHA3B	3884	NP_002270	NM_002279	4D	207787_at
arylsulfatase D	ARSD	414	NP_033667	NM_009589	4D	207784_at
presenilin 1 (Alzheimer disease 3)	PSEN1	5663	NP_015558	NM_007319	4D	207782_s_at
early growth response 4	EGR4	1961	NP_001956	NM_001965	4D	207768_at
DKFZP586A0522 protein	DKFZP586A052 2	25840	NP_054752	NM_014033	4D	207761_s_at
zinc finger protein 304	ZNF304	57343	NP_065708	NM_020657	4D	207753_at
NCKassociated protein 1	NCKAPI	10787	NP_038464	NM_013436	4D	207738_s_at
mutY homolog (E. coli)	HYTUM	4595	NP_036354	NM_012222	4D	207727_s_at
killer cell lectinlike receptor subfamily C, membe	KLRC3	3823	NP_002252	NM_002261	4D	207723_s_at
histidine triad nucleotide binding protein 1	HINTI	3094	NP_005331	NM_005340	4D	207721_x_at
KIAA0980 protein	KIAA0980	22981	NP_079452	NM_025176	4D	207705_s_at
leukocyte immunoglobulinlike receptor, subfamily B	LILRB2	10288	NP_005865	NM_005874	4D	207697_x_at
inhibin, beta C	INHBC	3626	NP_005529	NM_005538	4D	207687_at
one cut domain, family member 2	ONECUT2	9480	NP_004843	NM_004852	4D	207676_at

oxysterol binding proteinlike 1A	OSBPLIA	1114876	NP_060500	NM_018030	4D	208158_s_at
epiplakin I	EPPK1	83481	NP_112598	NM_031308	4D	208156_x_at
FAT tumor suppressor homolog 2 (Drosophila)	FAT2	2196	NP_001438	NM_001447	4D	208153_s_at
DEAD/H (AspGluAlaAsp/His) box polypeptide 11 (CHL1	DDX11	1663	NP_085911	NM_030653	40	208149_x_at
hypothetical protein MGC4293	MGC4293	83475	NP_112594	NM_031304	4D	208141_s_at
gastrin	GAS	2520	NP_000796	NM_000805	4D	208138_at
LASIlike (S. cerevisiae)	LASIL	81887	NP_112483	NM_031206	4D	208117_s_at
chromosome 9 open reading frame 74	C9orf74	81605	NP_112176	NM_030914	4D	208101_s_at
hypothetical protein MGC10471	MGC10471	81576	NP_110445	NM_030818	4D	208094_s_at
nudE nuclear distribution gene E homolog like 1 (A	NDELI	81565	NP_110435	NM_030808	4D	208093_s_at
family with sequence similarity 49, member A	FAM49A	81553	NP_110424	NM_030797	4D	208092_s_at
tudor domain containing 3	TDRD3	81550	NP_110421	NM_030794	4D	208089_s_at
dystrophin (muscular dystrophy, Duchenne and Becke	DMD	1756	NP_004005	NM_004014	4D	208086_s_at
integrin, beta 6	ITGB6	3694	NP_000879	NM_000888	4D	208084_at
integrin, beta 6	ITGB6	3694	NP_000879	NM_000888	4D	208083_s_at
				NM_015877	4D	208061_at
guanylate cyclase 2F, retinal	GUCY2F	2986	NP_001513	NM_001522	4D	208053_at
NGFIA binding protein 1 (EGR1 binding protein 1)	NABI	4664	NP_005957	NM_005966	4D	208047_s_at
				NM_013303	4D	208042_at
regulatory factor X, 2 (influences HLA class II ex	RFX2	5990	NP_000626	NM_000635	4D	208031_s_at
glutathione peroxidase 5 (epididymal androgenrelat	GPX5	2880	NP_003987	NM_003996	4D	208028_s_at
Ctype lectin domain family 4, member M	CLEC4M	10332	NP_055072	NM_014257	4D	207995_s_at
acrosomal vesicle protein 1	ACRV1	56	NP_064498	NM_020113	4D	207991_x_at
acrosomal vesicle protein l	ACRV1	56	NP_064499	NM_020114	4D	207990_x_at
CD8 antigen, beta polypeptide 1 (p37)	CD8B1	926	NP_004922	NM_004931	4D	207979_s_at
acrosomal vesicle protein 1	ACRV1	56	NP_064495	NM_020110	4D	207973_x_at
acrosomal vesicle protein 1	ACRV1	56	NP_064494	NM_020109	4D	207969_x_at
golgi apparatus protein 1	GLG1	2734	NP_036333	NM_012201	4D	207966_s_at
casein kinase I, delta	CSNKID	1453	NP_001884	NM_001893	4D	207945_s_at
6phosphofructo2kinase/fructose2,6biphosphatase	PFKFB2	5208	NP_006203	NM_006212	4D	207931_s_at

interferon, alpha 6	IFNA6	3443	NP_066282	NM_021002	4D	208548_at
histone 1, H2bh	HIST1H2BH	8345	NP_003515	NM_003524	4D	208546_x_at
histone 1, H2be	HIST1H2BE	8344	NP_003514	NM_003523	4D	208527_x_at
basic transcription factor 3	BTF3	689	NP_001198	NM_001207	4D	208517_x_at
olfactory receptor, family 2, subfamily J, member	OR2J2	26707	NP_112167	NM_030905	4D	208508_s_at
histone 1, H3f	HIST1H3F	8968	NP_066298	NM_021018	4D	208506_at
amylase, alpha 1A; salivary	AMYIA	276	NP_004029	NM_004038	4D	208498_s_at
histone 1, H3g	HIST1H3G	8355	NP_003525	NM_003534	4D	208496_x_at
histone 1, H2bf	HIST1H2BF	8343	NP_003513	NM_003522	4D	208490_x_at
RAB3D, member RAS oncogene family	RAB3D	9545	NP_004274	NM_004283	4D	208466_at
gap junction protein, alpha 7, 45kDa (connexin 45)	GJA7	10052	NP_005488	NM_005497	4D	208460_at
poliovirus receptorrelated 1 (herpesvirus entry me	PVRL1	5818	NP_002846	NM_002855	4D	208455_at
lectin, galactosidebinding, soluble, 2 (galectin 2	LGALS2	3957	NP_006489	NM_006498	4D	208450_at
bromodomain adjacent to zinc finger domain, 1B	BAZIB	9031	NP_075381	NM_023005	4D	208445_s_at
transporter 2, ATPbinding cassette, subfamily B (M	TAP2	6891	NP_061313	NM_018833	4D	208428_at
cytokine induced apoptosis inhibitor 1	CIAPINI	57019	NP_064709	NM_020313	4D	208424_s_at
TBPlike 1	TBPL1	9519	NP_004856	NM_004865	4D	208398_s_at
glucagonlike peptide 1 receptor	GLPIR	2740	NP_002053	NM_002062	4D	208391_s_at
solute carrier family 1 (glial high affinity gluta	SLC1A2	6506	NP_004162	NM_004171	4D	208389_s_at
capping protein (actin filament) muscle Zline, alp	CAPZAI	829	NP_006126	NM_006135	4D	208374_s_at
glutarylCoenzyme A dehydrogenase	GCDH	2639	NP_039663	NM_013976	4D	208369_s_at
protocadherin 11 Xlinked	PCDH11X	27328	NP_055337	NM_014522	4D	208366_at
PTPN13like, Ylinked	PRY	9081	NP_004667	NM_004676	4D	208332_at
A kinase (PRKA) anchor protein 13	AKAP13	11214	NP_006729	NM_006738	4D	208324_at
minor histocompatibility antigen HB1	HB1	57824	NP_067005	NM_021182	4D	208302_at
olfactory receptor, family 7, subfamily A, member	OR7A5	26659	NP_059976	NM_017506	4D	208285_at
ephrinA2	EFNA2	1943	NP_001396	NM_001405	4D	208256_at
TDPglucose 4,6dehydratase	TGDS	23483	NP_055120	NM_014305	4D	208249_s_at
				NM_012185	4D	208239_at
activin A receptor, type 1B	ACVRIB	91	NP_064732	NM_020327	4D	208223_s_at
interferon, alpha 14	IFNA14	3448	NP_002163	NM_002172	4D	208182_x_at
histone 1, H4h	HIST1H4H	8365	NP_003534	NM_003543	4D	208180_s_at
interleukin 9 receptor	IL9R	3581	NP_002177	NM_002186	4D	208164_s_at

proteasome (prosome, macropain) subunit, beta type	PSMB6	5694	NP_002789	NM_002798	4D	208827_at
histidine triad nucleotide binding protein 1	HINTI	3094	NP_005331	NM_005340	4D	208826_x_at
ribosomal protein L23a	RPL23A	6147	NP_000975	NM_000984	4D	208825_x_at
clusterin (complement lysis inhibitor, SP40,40, su	CLU	1191	NP_976084	NM_203339	4D	208791_at
polymerase I and transcript release factor	PTRF	284119	NP_036364	NM_012232	4D	208789_at
mitochondrial ribosomal protein L3	MRPL3	11222	NP_009139	NM_007208	4D	208787_at
microtubuleassociated protein 1 light chain 3 beta	MAP1LC3B	81631	NP_073729	NM_022818	4D	208786_s_at
microtubuleassociated protein 1 light chain 3 beta	MAPILC3B	81631	NP_073729	NM_022818	4D	208785_s_at
membrane cofactor protein (CD46, trophoblastlympho	MCP	4179	NP_758865	NM_172355	4D	208783_s_at
follistatinlike 1	FSTL1	11167	NP_009016	NM_007085	4D	208782_at
discoidin domain receptor family, member 1	DDRI	780	NP_054699	NM_013993	4D	208779_x_at
ribosomal protein L22	RPL22	6146	NP_000974	NM_000983	4D	208768_x_at
heterogeneous nuclear ribonucleoprotein R	HNRPR	10236	NP_005817	NM_005826	4D	208766_s_at
delta sleep inducing peptide, immunoreactor	DSIPI	1831	NP_004080	NM_004089	4D	208763_s_at
A 1 P Synthase, H+ transporting, mitochondrial F0 co	AIFOL	10032	NF_00040/	[NIV]_000+/0	ŧ	200/40_x_at
sin3associated polypeptide, 18kDa	SAP18	10284	NP_005861	NM_005870	100	208742 s_at
sin3associated polypeptide, 18kDa	SAP18	10284	NP_005861	NM_005870	4D	208740_at
RAB2, member RAS oncogene family	RAB2	5862	NP_002856	NM_002865	4D	208734_x_at
cyclin D1 (PRAD1: parathyroid adenomatosis 1)	CCND1	595	NP_444284	NM_053056	4D	208711_s_at
ribosomal protein L39	RPL39	6170	NP_000991	NM_001000	4D	208695_s_at
peroxiredoxin 1	PRDX1	5052	NP_002565	NM_002574	4D	208680_at
proliferationassociated 2G4, 38kDa	PA2G4	5036	NP_006182	NM_006191	4D	208676_s_at
dolichyldiphosphooligosaccharideprotein glycosyltr	DDOST	1650	NP_005207	NM_005216	4D	208674_x_at
ribosomal protein S14	RPS14	6208	NP_005608	NM_005617	4D	208646_at
thioredoxin domain containing 7 (protein disulfide	TXNDC7	10130	NP_005733	NM_005742	4D	208638_at
leukocyte immunoglobulinlike receptor, subfamily B	LILRB6	79168	NP_077294	NM_024318	4D	208594_x_at
corticotropin releasing hormone receptor 1	CRHR1	1394	NP_004373	NM_004382	4D	208593_x_at
H2B histone family, member S	H2BFS	54145	NP_059141	NM_017445	4D	208579_x_at
histone 1, H3c	HIST1H3C	8352	NP_003522	NM_003531	4D	208577_at
acidic (leucinerich) nuclear phosphoprotein 32 fam	ANP32D	23519	NP_036536	NM_012404	4D	208571_at

synaptotagmin binding, cytoplasmic RNA interacting	SYNCRIP	10492	NP_006363	NM_006372	4D	209025_s_at
KIAA0652 gene product	KIAA0652	9776	NP_055556	NM_014741	4D	209021_x_at
PTEN induced putative kinase 1	PINK1	65018	NP_115785	NM_032409	4D	209019_s_at
DnaJ (Hsp40) homolog, subfamily B, member 6	DNAJB6	10049	NP_490647	NM_058246	4D	209015_s_at
keratin 8	KRT8	3856	NP_002264	NM_002273	4D	209008_x_at
NPD014 protein	NPD014	57035	NP_996918	NM_207035	4D	209006_s_at
Fbox and leucinerich repeat protein 5	FBXL5	26234	NP_036293	NM_012161	4D	209004_s_at
anaphase promoting complex subunit 13	ANAPC13	25847	NP_056206	NM_015391	4D	209001_s_at
transcription factor 12 (HTF4, helixloophelix tran	TCF12	6938	NP_996921	NM_207038	4D	208986_at
uroporphyrinogen decarboxylase	UROD	7389	NP_000365	NM_000374	4D	208970_s_at
NADH dehydrogenase (ubiquinone) I alpha subcomplex	NDUFA9	4704	NP_004993	NM_005002	4D	208969_at
Kruppellike factor 6	KLF6	1316	NP_001291	NM_001300	4D	208961_s_at
Kruppellike factor 6	KLF6	1316	NP_001291	NM_001300	4D	208960_s_at
thioredoxin domain containing 4 (endoplasmic retic	TXNDC4	23071	NP_055866	NM_015051	4D	20 8957_a t
protein phosphatase 4 (formerly X), catalytic subu	PPP4C	5531	NP_002711	NM_002720	4D	208932_at
ring finger protein 11	RNFII	26994	NP_055187	NM_014372	4D	208924_at
NAD kinase	FLJ13052	65220	NP_075394	NM_023018	4D	208917_x_at
golgi associated, gamma adaptin ear containing, AR	GGA2	23062	NP_619581	NM_138640	4D	208914_at
golgi associated, gamma adaptin ear containing, AR	GGA2	23062	NP_055859	NM_015044	4D	208913_at
ribosomal protein S28	RPS28	6234	NP_001022	NM_001031	4D	208903_at
DEAD (AspGluAlaAsp) box polypeptide 18	DDX18	8886	NP_006764	NM_006773	4D	208896_at
DEAD (AspGluAlaAsp) box polypeptide 18	DDX18	8886	NP_006764	NM_006773	4D	208895_s_at
dual specificity phosphatase 6	DUSP6	1848	NP_001937	NM_001946	4D	208892_s_at
chromosome 5 open reading frame 18	C5orf18	7905	NP_005660	NM_005669	4D	208873_s_at
thioredoxin	TXN	7295	NP_003320	NM_003329	4D	208864_s_at
splicing factor, arginine/serinerich 1 (splicing f	SFRS1	6426	NP_008855	NM_006924	4D	208863_s_at
cisplatin resistanceassociated overexpressed prote	CROP	51747	NP_006098	NM_006107	4D	208835_s_at
ribosomal protein L23a	RPL23A	6147	NP_000975	NM_000984	4D	208834_x_at
ataxin 10	ATXN10	25814	NP_037368	NM_013236	4D	208833_s_at
polymerase (DNA directed), epsilon 3 (p17 subunit)	POLE3	54107	NP_059139	NM_017443	4D	208828_at

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carbonic anhydrase II	CA2	760	NP_000058	NM_000067	4D	209301_at
glutaredoxin (thioltransferase)	GLRX	2745	NP_002055	NM_002064	4D	209276 s at
vacuolar protein sorting 45A (yeast)	VPS45A	11311	NP_009190	NM_007259	4D	209268_at
methyltransferase like 3	METTL3	56339	NP_062826	NM_019852	4D	209265_s_at
tubulin alpha 6	TUBA6	84790	NP_116093	NM_032704	4D	209251_x_at
chloride channel 7	CLCN7	1186	NP_001278	NM_001287	4D	209235_at
C2f protein	C2F	10436	NP_006322	NM_006331	4D	209233_at
WD repeat domain 45	WDR45	111152	NP_009006	NM_007075	4D	209217_s_at
Kruppellike factor 5 (intestinal)	KLF5	688	NP_001721	NM_001730	4b	209211_at
downregulator of transcription 1, TBPbinding (nega	DRI	1810	NP_001929	NM_001938	40	209188_x_at
downregulator of transcription 1, TBPbinding (nega	DRI	1810	NP_001929	NM_001938	1 40	20918/_at
insulin receptor substrate 2	IRS2	8660	NP_003740	NM_003749	4D	209185_s_at
insulin receptor substrate 2	IRS2	8660	NP_003740	NM_003749	4D	209184_s_at
leukocyte receptor cluster (LRC) member 4	LENG4	79143	NP_077274	NM_024298	4D	
apoptosis antagonizing transcription factor	AATF	26574	NP_036270	NM_012138	4D	209165_at
pleckstrin homology, Sec7 and coiledcoil domains	PSCD2	9266	NP_059431	NM_01/45/	40	209138_S_at
transcription factor 3 (E2A immunoglobulin enhance	TCF3	6929	NP_003191	NM_003200	1	209152_s_at
transmembrane 9 superfamily member 1	TM9SF1	10548	NP_006396	NM_006405	140	209149_s_at
corebinding factor, runt domain, alpha subunit 2;	CBFA2T2	9139	NP_005084	NM_005093	4D	209144_s_at
synaptosomalassociated protein, 23kDa	SNAP23	8773	NP_003816	NM_003825	4D	209130_at
squamous cell carcinoma antigen recognised by T	SART3	9733	NP_055521	NM_014706	4D	209128_s_at
tubulin, alpha 3	TUBA3	7846	NP_006000	NM_006009	4D	209118_s_at
tetraspan 1	TSPANI	10103	NP_005718	NM_005727	4D	209114_at
nuclear receptor coactivator 1	NCOAI	8648	NP_671756	NM_147223	4D	209106_at
interferonrelated developmental regulator 2	IFRD2	7866	NP_006755	NM_006764	4D	209100_at
jagged 1 (Alagille syndrome)	JAG1	182	NP_000205	NM_000214	4D	209099 x_at
RAB5A, member RAS oncogene family	RAB5A	5868	NP_004153	NM_004162	4D	209089_at
collagen, type XVIII, alpha 1	COL18A1	80781	NP_085059	NM_030582	4D	209081_s_at
H3 histone, family 3B (H3.3B)	H3F3B	3021	NP_005315	NM_005324	4D	209069_s_at
ubiquinolcytochrome c reductase binding protein	UQCRB	7381	NP_006285	NM_006294	4D	209065_at
protein kinase C binding protein 1	PRKCBP1	23613	NP_898869	NM_183048	4D	209048_s_at

S100 calcium binding protein, beta (neural)	S100B	6285	NP_006263	NM_006272	4D	209686_at
carboxylesterase 2 (intestine, liver)	CES2	8824	NP_932327	NM_198061	4D	209667_at
CDC16 cell division cycle 16 homolog (S. cerevisia	CDC16	8881	NP_003894	NM_003903	4D	1
promyelocytic leukemia	PML	5371	NP_150241	NM_033238	4D	209640_at
regulator of Gprotein signalling 12	RGS12	6002	NP_937870	NM_198227	4D	209639_s_at
DNA segment on chromosome 4 (unique) 234 expressed	D4S234E	27065	NP_055207	NM_014392	4D	209569_x_at
RRS1 ribosome biogenesis regulator homolog (S. cer	RRS1	23212	NP_055984	NM_015169	45	20956/_at
hypothetical protein MGC11061	MGC11061	84272	NP_115688	NM_032312	4D	209551_at
exosome component 2	EXOSC2	23404	NP_055100	NM_014285	4D	209527_at
ash2 (absent, small, or homeotic)like (Drosophila)	ASH2L	9070	NP_004665	NM_004674	4D	209517_s_at
hydroxysteroid dehydrogenase like 2	HSDL2	84263	NP_115679	NM_032303	4D	209513_s_at
hydroxysteroid dehydrogenase like 2	HSDL2	84263	NP_115679	NM_032303	4D	209512_at
polymerase (RNA) II (DNA directed) polypeptide F	POLR2F	5435	NP_068809	NM_021974	4D	209511_at
RNA binding motif protein 30	RBM30	83759	NP_113680	NM_031492	4D	209497_s_at
chromosome 1 open reading frame 48	Clorf48	25936	NP_056286	NM_015471	4D	209484_s_at
ubiquitin specific protease 15	USP15	9958	NP_006304	NM_006313	4D	209475_at
TRAF family memberassociated NFKB activator	TANK	10010	NP_004171	NM_004180	4D	209451_at
Osialoglycoprotein endopeptidase	OSGEP	55644	NP_060277	NM_017807	4D	209450_at
phosphoribosyl pyrophosphate amidotransferase	PPAT	5471	NP_002694	NM_002703	4D	209434_s_at
alphamethylacylCoA racemase	AMACR	23600	NP_055139	NM_014324	4D	209426_s_at
chromosome 22 open reading frame 19	C22orf19	8563	NP_001002878	NM_001002878	4D	209418_s_at
transmembrane protein 1	TMEM1	7109	NP_003265	NM_003274	4D	209412_at
growth factor receptorbound protein 10	GRB10	2887	NP_001001550	NM_001001550	4D	209410_s_at
growth factor receptorbound protein 10	GRB10	2887	NP_001001549	NM_001001549	4D	209409_at
cytochrome b5	CYB5	1528	NP_001905	NM_001914	4D	209366_x_at
RAD50 homolog (S. cerevisiae)	RAD50	10111	NP_597816	NM_133482	4D	209349_at
MYC associated factor X	MAX	4149	NP_660088	NM_145113	4D	209331_s_at
hypothetical protein MGC2198	MGC2198	192286	NP_620175	NM_138820	4D	209328_x_at
polymerase (RNA) I polypeptide C, 30kDa	POLRIC	9533	NP_976035	NM_203290	4D	209317_at
BCL2/adenovirus E1B 19kDa interacting protein 2	BNIP2	663	NP_004321	NM_004330	4D	209308_s_at
growth arrest and DNAdamageinducible, beta	GADD45B	4616	NP_056490	NM_015675	4D	209305_s_at

oxidised low density lipoprotein (lectinlike) rece	OLRI	4973	NP_002534	NM_002543	4D	210004_at
suppressor of cytokine signaling 1	SOCS1	8651	NP_003736	NM_003745	4D	209999_x_at
zinc finger protein 268	ZNF268	10795	NP_003406	NM_003415	4D	209989_at
jumonji domain containing 2C	JMJD2C	23081	NP_055876	NM_015061	4D	209984_at
lipoprotein, Lp(a)	LPA	4018	NP_005568	NM_005577	4D	209978_s_at
JTV1 gene	JTV1	7965	NP_006294	NM_006303	4D	209971_x_at
melanoma antigen family A, 3	MAGEA3	4102	NP_005353	NM_005362	4D	209942_x_at
ATPase, Ca++ transporting, type 2C, member 1	ATP2C1	27032	NP_001001487	NM_001001487	4D	209934_s_at
CD300A antigen	CD300A	11314	NP_009192	NM_007261	4D	209933_s_at
	HIST1H2BD	3017	NP_619790	NM_138720	4D	209911_x_at
ataxia telangiectasia and Rad3 related	ATR	545	NP_001175	NM_001184	4D	209902_at
SEC31like 2 (S. cerevisiae)	SEC31L2	25956	NP_937781	NM_198138	4D	209889_at
Raslike without CAAX 1	RIT1	6016	NP_008843	NM_006912	4D	209882_at
linker for activation of T cells	LAT	27040	NP_055202	NM_014387	4D	209881_s_at
selectin P ligand	SELPLG	6404	NP_002997	NM_003006	4D	209880_s_at
synuclein, gamma (breast cancerspecific protein 1)	SNCG	6623	NP_003078	NM_003087	4D	209877_at
adrenergic, alpha2A, receptor	ADRA2A	150	NP_000672	NM_000681	4D	209869_at
latrophilin 3	LPHN3	23284	NP_056051	NM_015236	4D	209867_s_at
frequently rearranged in advanced Tcell lymphomas	FRAT2	23401	NP_036215	NM_012083	40	209864_at
tumor protein p73like	TP73L	8626	NP_003713	NM_003722	4D	209863_s_at
proteasome (prosome, macropain) activator subunit	PSME3	10197	NP_789839	NM_176863	4D	209853_s_at
CDC42 effector protein (Rho GTPase binding) 2	CDC42EP2	10435	NP_006770	NM_006779	4D	209850_s_at
leucine rich repeat neuronal 3	LRRN3	54674	NP_060804	NM_018334	4D	209841_s_at
leucine rich repeat neuronal 3	LRRN3	54674	NP_060804	NM_018334	4D	209840_s_at
palmitoylprotein thioesterase 2	PPT2	9374	NP_620312	NM_138934	4D	209826_at
jagged 2	JAG2	3714	NP_002217	NM_002226	4D	209784 s_at
CD24 antigen (small cell lung carcinoma cluster 4	CD24	934	NP_037362	NM_013230	4D	209772_s_at
itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	83737	NP_113671	NM_031483	4D	209744_x_at
itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	83737	NP_113671	NM_031483	4D	209743_s_at
Ctype lectin domain family 2, member B	CLEC2B	9976	NP_005118	NM_005127	4D	209732_at
solute carrier family 35 (UDPglucuronic acid/UDPNa	SLC35D1	23169	NP_055954	NM_015139	4D	209711_at
fatso	FTO	79068	XP_051200	XM_051200	4D	209702_at

membranespanning 4domains, subfamily A, member 3 (MS4A3	932	NP_006129	NM_006138	4D	210254_at
adenylosuccinate lyase	ADSL	158	NP_000017	NM_000026	4D	210250_x_at
cyclindependent kinase inhibitor 2D (p19, inhibits	CDKN2D	1032	NP_524145	NM_079421	4D	210240_s_at
protein tyrosine phosphatase, receptor type, f pol	PPFIAI	8500	NP_803172	NM_177423	4D	210236_at
interleukin 1 receptor accessory protein	ILIRAP	3556	NP_608273	NM_134470	4D	210233_at
transferrin receptor 2	TFR2	7036	NP_003218	NM_003227	4D	210215_at
syntaxin 11	STX11	8676	NP_003755	NM_003764	4D	210190_at
FK506 binding protein 1A, 12kDa	FKBPIA	2280	NP_000792	NM_000801	4D	210186_s_at
tolllike receptor 1	TLRI	7096	NP_003254	NM_003263	4D	210176_at
protein tyrosine phosphatase, receptor type, J	PTPRJ	5795	NP_002834	NM_002843	4D	210173_at
plateletactivating factor acetylhydrolase, isoform	PAFAH1B2	5049	NP_002563	NM_002572	4D	210160_at
leukocyte immunoglobulinlike receptor, subfamily B	LILRB4	11006	NP_006838	NM_006847	4D	210152_at
dCMP deaminase	DCTD	1635	NP_001912	NM_001921	4D	210137_s_at
leukotriene B4 receptor	LTB4R	1241	NP_858043	NM_181657	4D	210128_s_at
barrier to autointegration factor 1	BANFI	8815	NP_003851	NM_003860	4D	210125_s_at
potassium inwardlyrectifying channel, subfamily J,	KCNJ15	3772	NP_002234	NM_002243	4D	210119_at
sperm associated antigen 1	SPAGI	6674	NP_003105	NM_003114	4D	210117_at
inversin	INVS	27130	NP_899068	NM_183245	4D	210114_at
nucleolar protein 7, 27kDa	NOL7	51406	NP_057251	NM_016167	4D	210097_s_at
par3 partitioning defective 3 homolog (C. elegans)	PARD3	56288	NP_062565	NM_019619	4D	210094_s_at
magonashi homolog, proliferationassociated (Drosop	MAGOH	4116	NP_002361	NM_002370	4D	210093_s_at
magonashi homolog, proliferationassociated (Drosop	MAGOH	4116	NP_002361	NM_002370	4D	210092_at
myosin, light polypeptide 4, alkali; atrial, embry	MYL4	4635	NP_001002841	NM_001002841	4D	210088 x at
ribosomal protein L5	RPL5	6125	NP_000960	NM_000969	4D	210034_s_at
APEX nuclease (multifunctional DNA repair enzyme)	APEX1	328	NP_542380	NM_080649	4D	210027_s_at
polycomb group ring finger 1	PCGF1	84759	NP_116062	NM_032673	4D	210023_s_at
mucosa associated lymphoid tissue lymphoma translo	MALTI	10892	NP_006776	NM_006785	4D	210017_at
hemopexin	НРХ	3263	NP_000604	NM_000613	4D	210013_at
solute carrier family 25 (mitochondrial carrier; c	SLC25A1	6576	NP_005975	NM_005984	4D	210010_s_at

phosphate regulating endopeptidase homolog, Xlinke	РНЕХ	5251	NP_000435	NM_000444	4D	210617_at
synaptogyrin I	SYNGR1	9145	NP_663791	NM_145738	4D	210613_s_at
myelin protein zerolike 1	MPZL1	9019	NP_003944	NM_003953	4D	210594_x_at
Ras proteinspecific guanine nucleotidereleasing fa	RASGRF1	5923	NP_002882	NM_002891	4D	210550_s_at
cancer/testis antigen 1B	CTAG1B	1485	NP_001318	NM_001327	4D	210546_x_at
fucosyltransferase 7 (alpha (1,3) fucosyltransfera	FUT7	2529	NP_004470	NM_004479	4D	210506_at
fibronectin 1	FNI	2335	NP_997639	NM_212474	4D	210495_x_at
proteasome (prosome, macropain) 26S subunit, nonAT	PSMD4	5710	NP_722544	NM_153822	4D	210459_at
chromosome 10 open reading frame 28	C10orf28	27291	NP_055287	NM_014472	4D	210455_at
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5L	10632	NP_006467	NM_006476	4D	210453_x_at
mitogenactivated protein kinase 14	MAPK14	1432	NP_001306	NM_001315	4D	210449_x_at
Sjogren syndrome antigen A2 (60kDa, ribonucleoprot	SSA2	6738	NP_004591	NM_004600	4D	210438_x_at
defensin, beta 1	DEFB1	1672	NP_005209	NM_005218	4D	210397_at
myosin, light polypeptide 4, alkali; atrial, embry	MYL4	4635	NP_002467	NM_002476	4D	210395_x_at
synovial sarcoma, X breakpoint 4	SSX4	6759	NP_005627	NM_005636	4D	210394_x_at
histone 1, H2bg	HIST1H2BG	8339	NP_003509	NM_003518	4D	210387_at
calcium channel, voltagedependent, alpha 1G subuni	CACNAIG	8913	NP_938198	NM_198384	4	210380_s_at
lymphocyte antigen 9	LY9	4063	NP_002339	NM_002348	4D	210370_s_at
Bcell CLL/lymphoma 11A (zinc finger protein)	BCLIIA	53335	NP_060484	NM_018014	4D	210347 s_at
myelin transcription factor 1	MYTI	4661	NP_004526	NM_004535	4D	210341_at
peptidylglycine alphaamidating monooxygenase COOHt	PAMCI	9182	NP_005438	NM_005447	4D	210335_at
BTB/POZ KELCH domain protein	ENC2	64410	NP_071925	NM_022480	4D	210307_s_at
l(3)mbtlike (Drosophila)	L3MBTL	26013	NP_056293	NM_015478	4D	210306_at
mab21like 2 (C. elegans)	MAB21L2	10586	NP_006430	NM_006439	4D	210303_at
xanthine dehydrogenase	XDH	7498	NP_000370	NM_000379	4D	210301_at
mitogenactivated protein kinase kinase kinase 7 in	MAP3K7IP2	23118	NP_055908	NM_015093	4D	210284_s_at
G proteincoupled receptor 18	GPR18	2841	NP_005283	NM_005292	4D	210279_at
cytochrome P450, family 2, subfamily B, polypeptid	CYP2B6	1555	NP_000758	NM_000767	4D	210272_at
neurogenic differentiation 2	NEUROD2	4761	NP_006151	NM_006160	4D	210271_at

Nacylsphingosine amidohydrolase (acid ceramidase)	ASAHI	427	NP_004306	NM_004315	4D	210980_s_at
FAST kinase	FASTK	10922	NP_148936	NM_033015	4D	210975_x_at
KIAA0669 gene product	KIAA0669	9819	NP_055594	NM_014779	4D	210954 s at
insulin promoter factor 1, homeodomain transcripti	IPF1	3651	NP_000200	NM_000209	4D	210937_s_at
ring finger protein (C3H2C3 type) 6	RNF6	6049	NP_898866	NM_183045	4D	210931_at
similar to FKSG30	LOC440915	440915	XP_496601	XM_496601	4D	210926_at
programmed cell death 10	PDCD10	11235	NP_009148	NM_007217	4D	210907_s_at
interleukin 13 receptor, alpha 1	IL13RA1	3597	NP_001551	NM_001560	4D	210904_s_at
TP53 activated protein 1	TP53AP1	11257	NP_009164	NM_007233	4D	210886_x_at
RAD17 homolog (S. pombe)	RAD17	5884	NP_579916	NM_133338	4D	210826_x_at
BTB and CNC homology 1, basic leucine zipper trans	BACHI	571	NP_001177	NM_001186	40	210818_s_at
DEAD (AspGluAlaAsp) box polypeptide 49	DDX49	54555	NP_061943	NM_019070	4D	210811_s_at
CD27binding (Siva) protein	SIVA	10572	NP_006418	NM_006427	4D	210792_x_at
carcinoembryonic antigenrelated cell adhesion mole	CEACAM3	1084	NP_001806	NM_001815	40	210/89_x_at
calcium/calmodulindependent protein kinase	CAMKK2	10645	NP_757380	NM_172226	; 4B	210/87_s_at
survival of motor neuron protein interacting prote	SIPI	8487	NP_003607	NM_003616	4D	210779_x_at
caspase 9, apoptosisrelated cysteine protease	CASP9	842	NP_001220	NM_001229	4D	210775_x_at
nuclear receptor coactivator 4	NCOA4	8031	NP_005428	NM_005437	4D	210774_s_at
formyl peptide receptorlike 1	FPRL1	2358	NP_001005738	NM_001005738	4D	210772_at
vyes1 Yamaguchi sarcoma viral related oncogene hom	LYN	4067	NP_002341	NM_002350	4D	210754_s_at
discs, large (Drosophila) homologassociated protei	DLGAP1	9229	NP_001003809	NM_001003809	4D	210750 s at
discoidin domain receptor family, member 1	DDR1	780	NP_001945	NM_001954	4D	210749_x_at
inositol 1,3,4triphosphate 5/6 kinase	ITPK1	3705	NP_055031	NM_014216	4Đ	210740_s_at
MYC associated factor X	MAX	4149	NP_002373	NM_002382	4D	210734_x_at
egflike module containing, mucinlike, hormone rece	EMR3	84658	NP_113960	NM_0323/1	1	210/24_at
WW domain containing oxidoreductase	WWOX	51741	NP_057457	NM_016373		210695 s_at
ubiquitin specific protease 15	USP15	9958	NP_006304	NM_006313	4D	210681_s_at
ribosomal protein L13a	RPL13A	23521	NP_036555	NM_012423	4D	210646_x_at
tetratricopeptide repeat domain 3	TTC3	7267	NP_003307	NM_003316	4D	210645_s_at

eukaryotic translation initiation factor 3, subuni	EIF3S9	8662	NP_003/42	NM_003/31	40	211201_8_at
KIAA1539	KIAA1539	80256	NP_079458	NM_025182	1	
synovial sarcoma, X breakpoint 4	SSX4	6759	NP_005627	NM_005636	4D	211425_x_at
peptidyl arginine deiminase, type IV	PADI4	23569	NP_036519	NM_012387	4D	211413_s_at
Fc fragment of IgG, low affinity IIc, receptor for	FCGR2C	9103	NP_001005412	NM_001005412	4D	211396_at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_982255	NM_203431	4D	211378_x_at
interleukin 1 receptor, type II	IL1R2	7850	NP_004624	NM_004633	4D	211372_s_at
leptin receptor	LEPR	3953	NP_002294	NM_002303	4D	211355_x_at
hemochromatosis	HFE	3077	NP_000401	NM_000410	4D	211330_s_at
albumin	ALB	213	NP_000468	NM_000477	4D	211298_s_at
ubiquitin protein ligase E3A (human papilloma viru	UBE3A	7337	NP_570853	NM_130838	4D	211285_s_at
nuclear transcription factor Y, gamma	NFYC	4802	NP_055038	NM_014223	4D	211251_x_at
SH3domain binding protein 2	SH3BP2	6452	NP_003014	NM_003023	4D	211250 s_at
cytochrome P450, family 4, subfamily A, polypeptid	CYP4A11	1579	NP_000769	NM_000778	40	211231_x_at
tumor necrosis factor receptor superfamily, imember	TNFRSF10C	8794	NP_003832	NM_003841	4D	211163_s_at
actinin, alpha l	ACTNI	87	NP_001093	NM_001102	4D	211160_x_at
NGFIA binding protein 1 (EGR1 binding protein 1)	NABI	4664	NP_005957	NM_005966	4D	211139 s_at
leukocyte immunoglobulinlike receptor, subfamily	LILRB3	11025	NP_006855	NM_006864	4D	211135_x_at
ectodysplasin A	EDA	1896	NP_001005611	NM_001005611	4D	211128_at
survival of motor neuron protein interacting prote	SIP1	8487	NP_003607	NM_003616	4D	2111115_x_at
survival of motor neuron protein interacting prote	SIPI	8487	NP_003607	NM_003616	4D	2111114_x_at
leukocyte immunoglobulinlike receptor, subfamily A	LILRA2	11027	NP_006857	NM_006866	4D	211102_s_at
preBcell leukemia transcription factor 2	PBX2	5089	NP_002577	NM_002586	4D	211097 s_at
diazepam binding inhibitor (GABA receptor modulato	DBI	1622	NP_065438	NM_020548	4D	211070_x_at
exostoses (multiple)like 3	EXTL3	2137	NP_001431	NM_001440	4D	211051_s_at
peroxisomal biogenesis factor 7	PEX7	5191	NP_000279	NM_000288	4D	211033_s_at
aldehyde dehydrogenase 3 family, member B1	ALDH3B1	221	NP_000685	NM_000694	4D	211004_s_at
growth factor receptorbound protein 10	GRB10	2887	NP_001001555	NM_001001555	4D	210999_s_at
tripartite motificontaining 23	TRIM23	373	NP_150231	NM_033228	4D	210995_s_at

cyclin E2	CCNE2	9134	NP_477097	NM_057749	4D	211814_s_at
potassium inwardlyrectifying channel, subfamily J,	KCNJ15	3772	NP_733933	NM_170737	4D	211806_s_at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	4D	211765_x_at
ubiquitinconjugating enzyme E2B (RAD6 homolog)	UBE2B	7320	NP_003328	NM_003337	4D	211763_s_at
tubulin alpha 6	TUBA6	84790	NP_116093	NM_032704	4D	211750_x_at
elastase 3A, pancreatic (protease E)	ELA3A	10136	NP_005738	NM_005747	4D	211738_x_at
Fc fragment of IgE, high affinity I, receptor for;	FCERIA	2205	NP_001992	NM_002001	4D	211734_s_at
synovial sarcoma, X breakpoint 3	SSX3	10214	NP_783642	NM_175711	4D	211731_x_at
BH3 interacting domain death agonist	BID	637	NP_932071	NM_197967	4D	211725_s_at
phosphatase and tensin homolog (mutated in multipl	PTEN	5728	NP_000305	NM_000314	4D	211711_s_at
IQ motif containing B1	ІQСВ1	9657	NP_055457	NM_014642	4D	211707_s_at
RNA binding motif protein 13	RBM13	84549	NP_115898	NM_032509	4D	211686_s_at
interferon gamma receptor 1	IFNGRI	3459	NP_000407	NM_000416	4D	211676_s_at
mole	CEACAIMO	7000	INI _00Z+/+	NIVI_002763	ŧ	£11007_at
iammin, ocia i	CEACAME	7680	ND 002202	NM 002483	4D	
laminin hata l	I AMBI	3917	NP 002282	NM 002291	4D	
interlenkin 13 recentor slahe 1	II_13R A1	3597	- 11	NM 001560	4D	211612 s at
mitochondrial ribosomal protein L9	MRPL9	65005	NP_113608	NM_031420	4D	211594_s_at
membrane cofactor protein (CD46, trophoblastlympho	MCP	4179	NP_758866	NM_172356	4D	211574_s_at
brain and reproductive organexpressed (TNFRSF1A mo	BRE	9577	NP_004890	NM_004899	4D	211566_x_at
mitogenactivated protein kinase 14	MAPK14	1432	NP_001306	NM_001315	4D	211561_x_at
guanylate cyclase 1, soluble, beta 3	GUCY1B3	2983	NP_000848	NM_000857	4D	211555_s_at
hydroxyprostaglandin dehydrogenase 15(NAD)	HPGD	3248	NP_000851	NM_000860	4D	211548_s_at
ribosomal protein S10	RPS10	6204	NP_001005	NM_001014	4D	211542_x_at
heat shock 70kDa protein 2	HSPA2	3306	NP_068814	NM_021979	4D	211538_s_at
protein tyrosine phosphatase, receptor type, N pol	PTPRN2	5799	NP_002838	NM_002847	4D	211534_x_at
prolinerich protein BstNI subfamily 1	PRB1	5542	NP_955386	NM_199354	4D	211531_x_at
HLAG histocompatibility antigen, class I, G	HLAG	3135	NP_002118	NM_002127	4D	211530_x_at
pleckstrin homology, Sec7 and coiledcoil domains	PSCD4	27128	NP_037517	NM_013385	4D	211521_s_at
myotubularin related protein 3	MTMR3	8897	NP_694691	NM_153051	4D	211507_s_at

subunit						
proteasome (prosome, macropain) activator	PSME4	23198	NP_055429	NM_014614	4D	212222_at
optic atrophy 1 (autosomal dominant)	OPAI	4976	NP_570845	NM_130832	4D	212213_x_at
UKFZF364G2022 protein	2 DKFZP364G202	23903	NF_050512	[NIVI_O1547]	ŧ	212207_41
	IFI I M3	10410	NP 066362	NM_021034	1 5	212203 x_at
chromosome 10 open reading frame 61	C10orf61	26123	NP_056446	NM_015631	40	at
expressed in Tcells and eosinophils in atopic derm	ETEA	23197	NP_055428	NM_014613	4D	212108_at
expressed in Tcells and eosinophils in atopic derm	ETEA	23197	NP_055428	NM_014613	4D	212106_at
karyopherin alpha 6 (importin alpha 7)	KPNA6	23633	NP_036448	NM_012316	4D	212101_at
Era Gproteinlike 1 (E. coli)	ERAL1	26284	NP_005693	NM_005702	4D	212087_s_at
caldesmon 1	CALD1	800	NP_149130	NM_033139	4D	212077_at
myeloid/lymphoid or mixedlineage leukemia (trithor	MLL	4297	NP_005924	NM_005933	4D	212076_at
transient receptor potential cation channel, subfa	TRPC4AP	26133	NP_056453	NM_015638	4D	212059_s_at
ring finger protein 167	RNF167	26001	NP_056343	NM_015528	4D	212047_s_at
exocyst complex component 7	EXOC7	23265	NP_056034	NM_015219	4D	212034_s_at
antigen identified by monoclonal antibody Ki67	MKI67	4288	NP_002408	NM_002417	4D	212021_s_at
gro	CD##	700	MI_001001020	MM_001001330		# 1 1 0 4 tt
(DAA) (1	CDAA	020	NID 001001200	NM 001001390		212014 v at
H3 histone, family 3B (H3 3B)	H3F3B	3021	NP 005315	NM 005324	4D	211999 at
major histocompatibility complex, class II, DP alp	HLADPAI	3113	NP_291032	NM_033554	4D	211990_at
collagen, type IV, alpha 1	COL4A1	1282	NP_001836	NM_001845	4D	211981_at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	4D	211978_x_at
heat shock protein 90kDa alpha (cytosolic), class A member 1	HSP90AA1	3320	NP_005339	NM_005348	4D	211969_at
collagen, type IV, alpha 2	COL4A2	1284	NP_001837	NM_001846	4D	211964_at
plasminogen activator, urokinase receptor	PLAUR	5329	NP_001005377	NM_001005377	4D	211924_s_at
prostaglandin E receptor 3 (subtype EP3)	PTGER3	5733	NP_942007	NM_198714	4D	211909_x_at
fucosyltransferase 6 (alpha (1,3) fucosyltransfera	FUT6	2528	NP_000141	NM_000150	4D	211885_x_at
carcinoembryonic antigenrelated cell adhesion mole	CEACAMI	634	NP_001703	NM_001712	4D	211883_x_at
carcinoembryonic antigenrelated cell adhesion mole	CEACAM7	1087	NP_008821	NM_006890	4D	211848_s_at
myeloid/lymphoid or mixedlineage leukemia (trithor	MLLT2	4299	NP_005926	NM_005935	4D	211826_s_at

FADsynthetase	PP591	80308	NP_079483	NM_025207	4D	212541_at
chromodomain helicase DNA binding protein Hike	СНДІГ	9557	NP_004275	NM_004284	4D	212539_at
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4D	212537_x_at
DNA segment, Chr 15, Wayne State University 75, ex	D15Wsu75e	27351	XP_039495	XM_039495	4D	212527_at
phosphodiesterase 8A	PDE8A	5151	NP_775658	NM_173456	4D	212522_at
chromosome 10 open reading frame 22	C10orf22	84890	NP_116193	NM_032804	4D	212500_at
KIAA0553 protein	KIAA0553	23131	NP_001002909	NM_001002909	4D	212487_at
DnaJ (Hsp40) homolog, subfamily C, member 13	DNAJC13	23317	NP_776184	NM_173823	4D	212467_at
fibronectin 1	FNI	2335	NP_002017	NM_002026	4D	212464_s_at
transcription factor binding to IGHM enhancer 3	TFE3	7030	NP_006512	NM_006521	4D	212457_at
general transcription factor IIIC, polypeptide 2,	GTF3C2	2976	NP_001512	NM_001521	4D	212429_s_at
septin 6	SEPT6	23157	NP_665798	NM_145799	4D	212415_at
U3 snoRNP protein 4 homolog	IMP4	92856	NP_219484	NM_033416	4D	212411_at
KIAA0082	KIAA0082	23070	NP_055865	NM_015050	4D	212380_at
phosphoribosylglycinamide formyltransferase, phosp	GART	2618	NP_000810	NM_000819	4D	2125/8_at
ATPase, Ca++ transporting, cardiac muscle, slow tw	ATP2A2	488	NP_001672	NM_001681	40	212361_s_at
KIAA0323	KIAA0323	23351	NP_056114	NM_015299	4D	212355_at
amine oxidase (flavin containing) domain 2	AOF2	23028	NP_055828	NM_015013	4D	212348_s_at
myosin ID	MYOID	4642	NP_056009	NM_015194	4D	212338_at
glucosamine (Nacetyl)6sulfatase (Sanfilippo diseas	GNS	2799	NP_002067	NM_002076	4D	212334_at
22 DKFZP564F0522 protein	DKFZP564F0522	25940	NP_056290	NM_015475	4D	212333_at
transcription factor Dp1	TFDP1	7027	NP_009042	NM_007111	4D	212330_at
KIAA0746 protein	KIAA0746	23231	NP_056002	NM_015187	4D	212314_at
proteasome (prosome, macropain) 26S subunit, nonAT	PSMD14	10213	NP_005796	NM_005805	4D	212296_at
lipin I	LPINI	23175	NP_663731	NM_145693	4D	212274_at
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4D	212270_x_at
serine (or cysteine) proteinase inhibitor, clade B	SERPINBI	1992	NP_109591	NM_030666	4D	212268_at
dystonin	DST	667	NP_899236	NM_183380	4D	212254_s_at
tubulin, alpha 1 (testis specific)	TUBAI	7277	NP_005991	NM_006000	4D	212242_at
Fbox protein 21	FBXO21	23014	NP_296373	NM_033624	4D	212229_s_at

Sjogren syndrome antigen A2 (60kDa, ribonucleoprot	SSA2	6738	NP_004591	NM_004600	4D	212839_s_at
polymerase (DNAdirected), delta 3, accessory subun	POLD3	10714	NP_006582	NM_006591	4D	212836_at
EGFlikedomain, multiple 5	EGFL5	1955	XP_376905	XM_376905	4D	212831_at
PAX transcription activation domain interacting pr	PAXIPIL	22976	NP_031375	NM_007349	4D	212825_at
DnaJ (Hsp40) homolog, subfamily B, member 5	DNAJB5	25822	NP_036398	NM_012266	4D	212817_at
KIAA0877 protein	KIAA0877	23333	XP_371891	XM_371891	4D	212792_at
ribosomal protein L13a	RPL13A	23521	NP_036555	NM_012423	4D	212790_x_at
HDCMA18P protein	HDCMA18P	51574	NP_057732	NM_016648	4D	212785_s_at
GTP_binding protein	GTP	92170	NP_612393	NM_138384	4D	212767_at
ankyrin repeat and sterile alpha motif domain cont	ANKSI	23294	NP_056060	NM_015245	4D	212747_at
pleckstrin homology domain containing, family E (w	PLEKHE1	23239	NP_919431	NM_194449	4D	212719_at
cMpl binding protein	LOC113251	113251	NP_443111	NM_052879	4D	212714_at
bicaudal D homolog 2 (Drosophila)	BICD2	23299	NP_001003800	NM_001003800	4D	212702_s_at
septin 10	SEPT10	151011	NP_653311	NM_144710	4D	212698_s_at
hypothetical protein LOC162427	LOC162427	162427	NP_835227	NM_178126	4D	212697_at
protein phosphatase 1H (PP2C domain containing)	HIMAD	57460	XP_350881	XM_350880	4D	212686_at
zinc finger protein 3 (A851)	ZNF3	7551	NP_116313	NM_032924	4D	212684_at
erythrocyte membrane protein band 4.11ike 3	EPB41L3	23136	NP_036439	NM_012307	4D	212681_at
KIAA0582	KIAA0582	23177	NP_055962	NM_015147	4D	212677_s_at
secreted protein, acidic, cysteinerich (osteonecti	SPARC	6678	NP_003109	NM_003118	4D	212667_at
TCDDinducible poly(ADPribose) polymerase	TIPARP	25976	NP_056323	NM_015508	4D	212665_at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	4D	212661_x_at
interleukin I receptor antagonist	ILIRN	3557	NP_776215	NM_173843	4D	212659_s_at
sorting nexin 4	SNX4	8723	NP_003785	NM_003794	4D	212652_s_at
DEAH (AspGluAlaHis) box polypeptide 29	DHX29	54505	NP_061903	NM_019030	4D	212648_at
protein tyrosine phosphataselike (proline instead	PTPLB	201562	NP_940684	NM_198402	4D	212640_at
(mouse)	(Z)	7444	NF_990/3/	[NIM_20002]	ŧ	212000_at
initochoridi lai ricosonilai proteni 331	OVI	20114	111 0000 TH	25020C MIN		212636 ot
protein tyrosine prospinatase, receptor type, C	MDDC31	100/10	ND 005821	NM 005830	40	212603 at
zinc finger protein 451	ZNF451	20030	NP_0563/0	NM 080022	3 5	21233/_at
signal transducer and activator of transcription 5	STAT5B	6777	NP_036580	NM_012448	4D	212550_at

		2653	NP 004474	NM 004483	4D	213129 s at
S TSPVIIka S	TSPYL5	85453	NP_277047	NM_033512	4D	213122_at
C9 DnaJ (Hsp40) homolog, subfamily C, member 9	DNAJC9	23234	NP_056005	NM_015190	4D	213092_x_at
A ribosomal protein L23a	RPL23A	6147	NP_000975	NM_000984	4D	213084_x_at
PLSC domain containing protein	LOC254531	254531	NP_705841	NM_153613	4D	213078_x_at
zinc finger, FYVE domain containing 26	ZFYVE26	23503	NP_056161	NM_015346	4D	213073_at
806 nuclear protein UKp68	FLJ11806	79882	NP_079100	NM_024824	4D	213063_at
Nterminal asparagine amidase	NTAN1	123803	NP_775745	NM_173474	4D	213062_at
microtubule associated serine/threonine kinase 3	MAST3	23031	XP_038150	XM_038150	4D	213045_at
	ODAG	57798	NP_066990	NM_021167	4D	213018_at
tripartite motifcontaining 37	TRIM37	4591	NP_056109	NM_015294	4D	213009_s_at
TL2 angiopoietinlike 2	ANGPTL2	23452	NP_036230	NM_012098	4D	213001_at
)870 KIAA0870 protein	KIAA0870	22898	NP_055772	NM_014957	4D	212974_at
ribose 5phosphate isomerase A (ribose 5phosphate e	RPIA	22934	NP_653164	NM_144563	4D	212973_at
hypermethylated in cancer 2	HIC2	23119	NP_055909	NM_015094	4D	212964_at
peptidylglycine alphaamidating monooxygenase	PAM	5066	NP_620177	NM_138822	4D	212958_x_at
21 polymerase (RNA) II (DNA directed) polypeptide I,	POLR21	5438	NP_006224	NM_006233	4D	212955_s_at
collagen, type VI, alpha l	COL6A1	1291	NP_001839	NM_001848	4D	212940_at
DKFZP564D172 hypothetical protein DKFZp564D172	DKFZF	83989	NP_114431	NM_032042	4D	212936_at
ribosomal protein L13	RPL13	6137	NP_000968	NM_000977	4D	212933_x_at
_4 TSPYlike 4	TSPYL4	23270	NP_067680	NM_021648	4D	212928_at
L1 SMC5 structural maintenance of chromosomes 51ike 1	SMC5L1	23137	NP_055925	NM_015110	4D	212926_at
21 chromosome 19 open reading frame 21	Cl9orf2	126353	NP_775752	NM_173481	4D	212925_at
2 SET and MYND domain containing 2	SMYD2	56950	NP_064582	NM_020197	4D	212922_s_at
cleavage stimulation factor, 3' preRNA, subunit 2,	CSTF2T	23283	NP_056050	NM_015235	4D	212905_at
SPH10 Mphase phosphoprotein 10 (U3 small nucleolar ribon	MPHOSPH10	10199	NP_005782	NM_005791	4D	212885_at
25 chromosome 21 open reading frame 25	C21orf25	25966	NP_950251	NM_199050	4D	212875_s_at
ubiquitin specific protease 49	USP49	9477	NP_004266	NM_004275	4D	212872_s_at
CDPdiacylglycerol synthase (phosphatidate cytidyly	CDS2	8760	NP_003809	NM_003818	4D	212862_at
)276 KIAA0276 protein	KIAA0276	23142	NP_055930	NM_015115	4D	212855_at

similar to hypothetical protein MGC13138	LOC146542	146542	NP_660314	NM_145271	4D	213527_s_at
cyclin E1	CCNE1	898	NP_476530	NM_057182	4D	213523_at
laminin, alpha 2 (merosin, congenital muscular dys	LAMA2	3908	NP_000417	NM_000426	4D	213519_s_at
coagulation factor II (thrombin) receptorlike 1	F2RL1	2150	NP_005233	NM_005242	4D	213506_at
acylCoenzyme A oxidase 1, palmitoyl	ACOX1	51	NP_009223	NM_007292	4D	213501_at
collagen, type II, alpha I (primary osteoarthritis	COL2A1	1280	NP_149162	NM_033150	4D	213492_at
KIAA0073 protein	KIAA0073	23398	NP_056157	NM_015342	4D	213483_at
vesicleassociated membrane protein 4	VAMP4	8674	NP_973723	NM_201994	4D	213480_at
malignant fibrous histiocytoma amplified sequence	MFHAS1	9258	NP_004216	NM_004225	4D	213457_at
similar to RUN and FYVE domaincontaining 2; Run an	LOC441022	441022	XP_496700	XM_496700	4D	213437_at
epimorphin	EPIM	2054	NP_919337	NM_194356	4D	213434_at
ribosomal protein S19	RPS19	6223	NP_001013	NM_001022	4D	213414_s_at
tight junction protein 3 (zona occludens 3)	TJP3	27134	NP_055243	NM_014428	4D	213412_at
hypothetical protein LOC126208	LOC126208	126208	NP_001002836	NM_001002836	4D	213402_at
chromosome 14 open reading frame 124	C14orf124	56948	NP_064580	NM_020195	4D	213398_s_at
DEAD/H (AspGluAlaAsp/His) box polypeptide 11 (CHL1	DDXII	1663	NP_004390	NM_004399	4D	213378_s_at
caspase 8, apoptosisrelated cysteine protease	CASP8	841	NP_001219	NM_001228	4D	213373_s_at
transmembrane and coiledcoil domains 1	TMCC1	23023	NP_055823	NM_015008	4D	213349_at
KIAA0495	KIAA0495	57212	NP_997189	NM_207306	4D	213340_s_at
vesicleassociated membrane protein I (synaptobrevi	VAMP1	6843	NP_055046	NM_014231	4D	213326_at
chromosome 6 open reading frame 162	C6orf162	57150	NP_065158	NM_020425	4D	213312_at
phosphoribosylformylglycinamidine synthase (FGAR a	PFAS	5198	NP_036525	NM_012393	4D	213302_at
zinc finger protein 248	ZNF248	57209	NP_066383	NM_021045	4D	213269_at
zinc finger protein 187	ZNF187	7741	NP_689949	NM_152736	4D	213218_at
small nuclear RNA activating complex, polypeptide	SNAPC5	10302	NP_006040	NM_006049	4D	213203_at
component of oligomeric golgi complex 7	COG7	91949	NP_705831	NM_153603	4D	213190_at
Sp3 transcription factor	SP3	6670	NP_003102	NM_003111	4D	213168_at
glycine cleavage system protein H (aminomethyl car	GCSH	2653	NP_004474	NM_004483	4D	213133_s_at

Nacylsphingosine amidohydrolase (acid	ASAHI	427	NP_004306	NM_004315	40	213902_at
adenine phosphoribosyltransferase	APRT	353	NP_000476	NM_000485	4D	213892_s_at
phosphatidylinositol glycan, class L	PIGL	9487	NP_004269	NM_004278	4D	213889_at
zinc finger, CSL domain containing 3	ZCSL3	120526	NP_859057	NM_181706	4D	213853_at
Similar to RIKEN cDNA 1810038N08 gene	MGC52022	375346	NP_940965	NM_198563	4D	213851_at
cytochrome c oxidase subunit VIIc	COX7C	1350	NP_001858	NM_001867	4D	213846_at
WD40 repeat protein Interacting with phosphoInosit	WIPI49	55062	NP_060453	NM_017983	4D	213836_s_at
GTP binding protein 3 (mitochondrial)	GTPBP3	84705	NP_116009	NM_032620	4D	213835_x_at
homeo box A11	HOXAII	3207	NP_005514	NM_005523	4D	213823_at
START domain containing 5	STARD5	80765	NP_085051	NM_030574	4D	213820_s_at
enhance	ICF3	0923	NF_003171	14141 000Z00	ŧ	213011_X_at
protein tyrosine phosphatase, receptor type, A	FIFNA	0000	NID 002 101	THOOOD TAIN	3 5	212011 2 24
CAP, adenylate cyclaseassociated protein 1 (yeast)	CAP1	1048/		NM_000841	3 5	
p					i	
Tay I (human Toell lankemia virus type I) hinding	TAXIRPI	8887	NP 006015	NM 006024	4D	213786 at
hypothetical protein FLJ43806	FLJ43806	399563	NP_963922	NM_201628	4D	213752_at
hypothetical protein LOC284352	LOC284352	284352	XP_113978	XM_113978	4D	213751_at
MAXlike protein X	MLX	6945	NP_937848	NM_198205	4D	213708_s_at
Rab geranylgeranyltransferase, beta subunit	RABGGTB	5876	NP_004573	NM_004582	4D	213704_at
ribosomal protein L35a	RPL35A	6165	NP_000987	NM_000996	4D	213687_s_at
NOL1/NOP2/Sun domain family, member 5	NSUN5	55695	NP_683759	NM_148956	4D	213670_x_at
septin 6	SEPT6	23157	NP_665799	NM_145800	4D	213666_at
zinc finger protein 75 (D8C6)	ZNF75	7626	NP_009062	NM_007131	4D	213659_at
similar to KIAA0363	LOC392666	392666	XP_374432	XM_374432	4D	213630_at
kinesin family member 3A	KIF3A	11127	NP_008985	NM_007054	4D	213623_at
NAD kinase	FLJ13052	65220	NP_075394	NM_023018	4D	213607_x_at
FUS interacting protein (serinearginine rich) 1	FUSIP1	10772	NP_473357	NM_054016	4D	213594_x_at
angiotensin II receptorlike 1	AGTRL1	187	NP_005152	NM_005161	4D	213592_at
ribosomal protein L14	RPL14	9045	NP_003964	NM_003973	4D	213588_x_at
E1A binding protein p300	EP300	2033	NP_001420	NM_001429	4D	213579_s_at
squalene epoxidase	SQLE	6713	NP_003120	NM_003129	4D	213577_at
sorting nexin 3	SNX3	8724	NP_003786	NM_003795	4D	213545_x_at
major histocompatibility complex, class II, DP alp	HLADPAI	3113	NP_291032	NM_033554	4D	213537_at

ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4D	214291_at
fatty acid binding protein 3, muscle and heart (ma	FABP3	2170	NP_004093	NM_004102	4D	214285_at
galanin	GAL	51083		NM_015973	4D	214240_at
protein (peptidylprolyl cis/trans isomerase) NIMAi	PIN4	5303	NP_006214	NM_006223	4D	214224_s_at
complement component 1, q subcomponent binding pro	CIQBP	708	NP_001203	NM_001212	4D	214214_s_at
solute carrier family 25 (mitochondrial carrier; p	SLC25A17	10478	NP_006349	NM_006358	4D	214210_at
DiGeorge syndrome critical region gene 2	DGCR2	9993	NP_005128	NM_005137	4D	214198_s_at
tripeptidyl peptidase I	TPP1	1200	NP_000382	NM_000391	4D	214195_at
nuclear factor (erythroidderived 2)like 1	NFE2L1	4779	NP_003195	NM_003204	4D	214179_s_at
fumarate hydratase	FH	2271	NP_000134	NM_000143	4D	214170_x_at
succinate dehydrogenase complex, subunit B, iron s	SDHB	6390	NP_002991	NM_003000	4D	214166_at
polymerase (RNA) II (DNA directed) polypeptide D	POLR2D	5433	NP_004796	NM_004805	4D	214144_at
ribosomal protein L24	RPL24	6152	NP_000977	NM_000986	4D	214143_x_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT13	25961	NP_056985	NM_015901	4D	214136_at
pericentriolar material 1	PCM1	5108	NP_006188	NM_006197	4D	214118_x_at
MYC associated factor X	MAX	4149	NP_660089	NM_145114	4D	214108_at
GDPmannose 4,6dehydratase	GMDS	2762	NP_001491	NM_001500	4D	214106_s_at
glutathione peroxidase 3 (plasma)	GPX3	2878	NP_002075	NM_002084	4D	214091_s_at
neutrophil cytosolic factor 1 (47kDa, chronic gran	NCF1	4687	NP_000256	NM_000265	4D	214084_x_at
singlestranded DNA binding protein 1	SSBP1	6742	NP_003134	NM_003143	4ם	214060_at
gelsolín (amyloidosis, Finnish type)	GSN	2934	NP_000168	NM_000177	4D	214040_s_at
chromosome X open reading frame 37	CXorf37	28952	NP_054727	NM_014008	4D	214037_s_at
CDC42 effector protein (Rho GTPase binding) 2	CDC42EP2	10435	NP_006770	NM_006779	4D	214014_at
hypothetical protein HSPC111	HSPC111	51491	NP_057475	NM_016391	4D	214011_s_at
ribosomal protein S20	RPS20	6224	NP_001014	NM_001023	4D	214003_x_at
myosin, light polypeptide 6, alkali, smooth muscle	MYL6	4637	NP_524148	NM_079424	4D	214002_at
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5S	27109	NP_001003803	NM_001003803	4D	213995_at
ADAMTSlike 3	ADAMTSL3	57188	NP_997400	NM_207517	4D	213974_at
ribosomal protein L29	RPL29	6159	NP_000983	NM_000992	4D	213969_x_at
formin binding protein 1	FNBP1	23048	NP_055848	NM_015033	4D	213940_s_at
ceramidase)						

menoral property sources		1111		NM 002228	45	214590 s at
vacualar protein sorting 52 (veget)	VPS52	6293	NP_072047	NM_022553	4D	214585_s_at
tumor necrosis factor receptor superfamily, member	TNFRSF21	27242	NP_055267	NM_014452	4D	214581_x_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_996735	NM_206853	4D	214543_x_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_996736	NM_206854	4D	214541_s_at
histone 1, H2bo	HIST1H2BO	8348	NP_003518	NM_003527	4D	214540_at
serine (or cysteine) proteinase inhibitor, clade B	SERPINB10	5273	NP_005015	NM_005024	4D	214539_at
				NM_005394	4D	214526_x_at
mutL homolog 3 (E. coli)	MLH3	27030	NP_055196	NM_014381	4D	214525_x_at
Fc fragment of IgG, high affinity la, receptor (CD	FCGRIA	2209	NP_000557	NM_000566	4D	214511_x_at
cAMP responsive element modulator	CREM	1390	NP_898883	NM_183060	4D	214508_x_at
exosome component 2	EXOSC2	23404	NP_055100	NM_014285	4D	214507_s_at
agouti signaling protein, nonagouti homolog (mouse	ASIP	434	NP_001663	NM_001672	4D	214498_at
gly						
sarcoglycan, delta (35kDa dystrophinassociated	SGCD	6444	NP_000328	NM_000337	4D	214492_at
serum amyloid A1	SAA1	6288	NP_000322	NM_000331	4D	214456_x_at
histone 1, H2bc	HIST1H2BC	8347	NP_003517	NM_003526	4D	214455_at
syntaxin 6	STX6	10228	NP_005810	NM_005819	4D	214441_at
Nacetyltransferase 1 (arylamine Nacetyltransferase	NATI	9	NP_000653	NM_000662	4D	214440_at
H2.0like homeo box 1 (Drosophila)	HLXI	3142	NP_068777	NM_021958	4D	214438_at
heat shock 70kDa protein 12A	HSPA12A	259217	XP_048898	XM_048898	4D	214434_at
nucleolar protein 1, 120kDa	NOLI	4839	NP_006161	NM_006170	4D	214427_at
methylCpG binding domain protein 2	MBD2	8932	NP_056647	NM_015832	4D	214397_at
arachidonate 5lipoxygenase	ALOX5	240	NP_000689	NM_000698	4D	214366_s_at
tropomyosin 3	TPM3	7170	NP_689476	NM_152263	4D	214365_at
acetylCoenzyme A carboxylase alpha	ACACA	31	NP_942131	NM_198834	4D	214358_at
dopa decarboxylase (aromatic Lamino acid decarboxy	DDC	1644	NP_000781	NM_000790	4D	214347_s_at
mitogenactivated protein kinase kinase kinase kina	MAP4K1	11184	NP_009112	NM_007181	4D	214339_s_at
nephroblastoma overexpressed gene	NOV	4856	NP_002505	NM_002514	4D	214321_at
septin 6	SEPT6	23157	NP_055944	NM_015129	4D	214298_x_at

transcription factor 3 (E2A immunoglobulin	TCF3	6929	NP_003191	NM_003200	4D	215260_s_at
eukaryotic translation initiation factor 3, subuni	EIF3S8	8663	NP_003743	NM_003752	4D	215230_x_at
keratin, hair, basic, 6 (monilethrix)	KRTHB6	3892	NP_002275	NM_002284	4D	215189_at
cadherinlike 22	CDH22	64405	NP_067071	NM_021248	4D	215181_at
NAD kinase	FLJ13052	65220	NP_075394	NM_023018	4D	215159_s_at
exosome component 8	EXOSC8	11340	NP_852480	NM_181503	4D	215136_s_at
similar to KIAA0752 protein	LOC389347	389347	XP_371783	XM_371783	4D	215133_s_at
KIAA0492 protein	KIAA0492	57238	XP_378914	XM_378914	4D	215109_at
NAD(P) dependent steroid dehydrogenaselike	NSDHL	50814	NP_057006	NM_015922	4D	215093_at
erythropoietin receptor	EPOR	2057	NP_000112	NM_000121	4D	215054_at
neurexin 3	NRXN3	9369	NP_004787	NM_004796	4D	215021_s_at
fasciculation and elongation protein zeta 2 (zygin	FEZ2	9637	NP_005093	NM_005102	4D	215000_s_at
glutamate receptor, ionotropic, kainate 5	GRIK5	2901	NP_002079	NM_002088	4D	214966_at
hypothetical protein MGC26885	MGC26885	124044	NP_689552	NM_152339	4D	214965_at
KIAA0527 protein	KIAA0527	26032	XP_171054	XM_171054	4D	214954_at
ATPase, Class II, type 9B	АТР9В	374868	NP_940933	NM_198531	4D	214934_at
SHC (Src homology 2 domain containing) transformin	SHC1	6464	NP_003020	NM_003029	å D	214853_s_at
Gprotein signalling modulator 3 (AGS3like, C. eleg	GPSM3	63940	NP_071390	NM_022107	4D	214847_s_at
docking protein 5	DOK5	55816	NP_060901	NM_018431	4D	214844_s_at
hypothetical protein from clone 24828	KIAA1747	57795	NP_066988	NM_021165	4D	214822_at
exportin 6	XPO6	23214	NP_055986	NM_015171	4D	214784_x_at
putative MAPK activating protein	MGC3794	261726	NP_690866	NM_152902	4D	214773_x_at
G2 protein	G2	25758	XP_039515	XM_039515	4D	214772_at
Sp1 transcription factor	SP1	6667	NP_612482	NM_138473	4D	214732_at
zinc finger protein 200	ZNF200	7752	NP_003445	NM_003454	4D	214706_at
InaDlike (Drosophila)	INADL	10207	NP_795353	NM_176878	4D	214705_at
hypothetical protein MGC14376	MGC14376	84981	NP_001001870	NM_001001870	4D	214696_at
neuropilin 2	NRP2	8828	NP_958436	NM_201279	4D	214632_at
glucosidase, alpha; neutral AB	GANAB	23193	NP_938148	NM_198334	4D	214626_s_at
CASP8 and FADDlike apoptosis regulator	CFLAR	8837	NP_003870	NM_003879	4D	214618_at
tetraspan 2	TSPAN2	10100	NP_005716	NM_005725	4D	214606_at
mornorog,						

NP_829844 23085 RAB6IP2 RAB6 interacting protein 2 NP_000422 4598 MVK mevalonate kinase (mevalonic aciduria) NP_000422 4598 MVK mevalonate kinase (mevalonic aciduria) NP_000422 4598 MVK mevalonate kinase (mevalonic aciduria) NP_000938 5558 PARS poly characteristic segregation increased 2 like 5 NP_000938 5558 PRIM2A primase, polypeptide 2A, 58kDa NP_000931 355 FAS Fas (TNF receptor superfamily, member 6) NP_00103806 445347 TARP TCR gamma alternate reading frame protein NP_00103806 445347 TARP TCR gamma alternate reading frame protein NP_0014143 4946 NAGLU Nacetylglucosaminidase, alpha (Sanfilippo disease NP_001621 4946 OAZI ornithine decarboxylase antizyme 1 NP_003215 10139 ARFRPI ADPribosylation factor related protein 1 NP_003215 10139 ARFRPI ADPribosylation factor related protein 1 NP_004854 9517 SPTICC2 serine palmitoyltransfe	806	215806_x_at 4D 215880_at 4D 215947_s_at 4D 215952_s_at 4D 215983_s_at 4D 215984_s_at 4D 216021_s_at 4D 216034_at 4D 216074_x_at 4D 216202_s_at 4D 216234_s_at 4D 216254_s_at 4D 216255_at 4D 216351_x_at 4D 216351_x_at 4D 216350_x_at 4D 216350_x_at 4D
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 0005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 0056520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA 002721 5566 PRKACA 775263 4929 NR4A2 001003828 29780 PARVB 000158 2918 GRM8 000158 2710 GK 065153 57135 DAZ4	828	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 0048540 9517 GLRA3 056053 23286 KIBRA 002721 5566 PRKACA 004854 9517 SPTLC2 004854 9517 SPTLC3 004854 9517 SPTKACA 0001003828 29780 PARVB 001003828 2918 GRM8 000158 2710 GK	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 003215 10139 ARFRP1 0048520 8001 GLRA3 056053 23286 KIBRA 004854 9517 SPTLC2 002721 5566 PRKACA 775263 4929 NR4A2 690612 355 FAS 001003828 29780 PARVB	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 0056520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA 002721 5566 PRKACA 775263 4929 NR4A2 001003828 29780 PARVB	828	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 006520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA 002721 5566 PRKACA 002721 5566 PRKACA 775263 4929 NR4A2 690612 355 FAS	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 003215 10139 ARFRP1 0048520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA 004854 9517 SPTLC2 0775263 4929 NR4A2	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 006520 8001 GLRA3 542778 129025 SUHWI 056053 23286 KIBRA 002721 5566 PRKACA	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 006520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA 004854 9517 SPTLC2	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 003215 10139 ARFRP1 006520 8001 GLRA3 542778 129025 SUHW1 056053 23286 KIBRA	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 003215 10139 ARFRP1 006520 8001 GLRA3 542778 129025 SUHW1	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E 006520 8001 GLRA3	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 690613 5558 PRIM2A 6905475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 003215 10139 ARFRP1	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1 005662 7993 D8S2298E	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668 004143 4946 OAZ1	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2 001003806 445347 TARP 000254 4669 NAGLU 116211 84908 FLJ14668	806	
23085 RAB6IP2 4598 MVK 5383 PMS2L5 5558 PRIM2A 355 FAS 10038 PARP2 445347 TARP 4669 NAGLU	806	
23085 RAB6IP2 4598 MVK 5383 PMS2L5 5558 PRIM2A 355 FAS 10038 PARP2 445347 TARP	806	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS 005475 10038 PARP2	NP	
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A 690613 355 FAS		215773_x_at 4D
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5 000938 5558 PRIM2A	NM 152874 NP 690613	215719_x_at 4D
829884 23085 RAB6IP2 000422 4598 MVK 777590 5383 PMS2L5		15708_s_at 4D
829884 23085 RAB6IP2 000422 4598 MVK		15667_x_at 4D
829884 23085 RAB6IP2		15649_s_at 4D
	NM_178040 NP_829884	15606_s_at 4D
NP_067684 11039 SMA4 SMA4		215599_at 4D
NP_057190 23256 SCFD1 sec1 family domain containing 1	11 1	215548_s_at 4D
NP_005210 1729 DIAPH1 diaphanous homolog 1 (Drosophila)		215541_s_at
NP_008980 11120 BTN2A1 butyrophilin, subfamily 2, member A1	NM_007049 NP_008980	215493_x_at
XP_043653 56271 BEXL1 brain expressed Xlinkedlike 1	XM_043653	215440_s_at
NP_038477 11176 BAZ2A bromodomain adjacent to zinc finger domain, 2A	NM_013449 NP_038477	215437_x_at 4D
NP_001001430 7139 TNNT2 troponin T2, cardiac	NM_001001430 NP_001001430	215389_s_at 4D
NP_742099 926 CD8B1 CD8 antigen, beta polypeptide 1 (p37)		15332_s_at 4D
NP_612147 154661 RPIB9 Rap2binding protein 9	1 1	15321_at 4D
NP_000330 6559 SLC12A3 solute carrier family 12 (sodium/chloride transpor		

GRIA3	2892	NP_015564	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		
25.			NM 007325	4D	217565 at
			XM_497371	4D	217499_x_at
PTEN	5728	NP_000305	NM_000314	4D	217492_s_at
COX7C	1350	NP_001858	NM_001867	4D	217491_x_at
IL6R	3570	NP_000556	NM_000565	4D	217489_s_at
LIMK2	3985	NP_005560	NM_005569	4D	217475_s_at
ADRB3	155	NP_000016	NM_000025	4D	217303_s_at
NBS1	4683	NP_002476	NM_002485	4D	217299 s_at
MLH3	27030	NP_055196	NM_014381	4D	217216_x_at
KIAA1078	23271	NP_982284	NM_203459	4D	217196_s_at
ZNF259	8882	NP_003895	NM_003904	4D	217186_at
GK	2710	NP_976325	NM_203391	4D	217167_x_at
MUC4	4585	NP_004523	NM_004532	4D	217109_at
ITCH	83737	NP_113671	NM_031483	4D	217094_s_at
ELK1	2002	NP_005220	NM_005229	4D	217039_x_at
LOC389177	389177	XP_371680	XM_371680	4D	217016_x_at
AZGP1	563	NP_001176	NM_001185	4D	217014_s_at
RIP	84268	NP_115684	NM_032308	4D	216962_at
FCGRIA	2209	NP_000557	NM_000566	Ð	216950_s_at
RSI	6247	NP_000321	NM_000330	4D	216937_s_at
DAZI	1617	NP_004072	NM_004081	4D	216922_x_at
PTPN12	5782	NP_002826	NM_002835	4D	216915_s_at
SOD2	6648	NP_000627	NM_000636	4D	216841_s_at
LOC442711	442711	XP_499457	XM_499457	4D	216824_at
GATM	2628	NP_001473	NM_001482	4D	216733_s_at
OR2B6	26212	NP_036499	NM_012367	4D	216522_at
PTMA	5757	NP_002814	NM_002823	4D	216515_x_at
MLLT10	8028	NP_004632	NM_004641	4D	216503_s_at
TMSL3	7117	NP_898870	NM_183049	4D	216438_s_at
ABCD1	215	NP_000024	NM_000033	4D	216418_at
	ABCDI TMSL3 TMSL3 MILT10 PTMA OR2B6 GATM LOC4427 SOD2 PTPN12 DAZ1 RS1 RS1 RCGR1A RIP AZGP1 LOC3891 ELK1 ITCH MUC4 GK ZNF259 KIAA107 MLH3 NBS1 ADRB3 LIMK2 IL6R		215 215 217 207117 28028 26212 26212 26212 442711 442711 6648 6247 6247 6247 6247 2209 389177 389177 2002 2155 23271 23271 23271 23271 23271 23271 23271 23271 2328 23271	000033 NP_000024 215 183049 NP_898870 7117 002823 NP_004632 8028 001482 NP_036499 26212 001482 NP_001473 2628 001482 NP_001473 2628 000636 NP_001473 2628 0004081 NP_000627 6648 000330 NP_0004072 1617 0005266 NP_000321 6247 000527 NP_001176 563 371680 NP_001176 563 371680 NP_005220 2002 031483 NP_005220 2002 371680 NP_003895 882 203391 NP_00520 2002 0044532 NP_003895 8882 203459 NP_0055196 27030 004885 NP_0055196 27030 005569 NP_0055196 27030 005569 NP_005560 3985 000556 NP_000556 3570	NM_000033 NP_000024 215 NM_183049 NP_898870 7117 NM_004641 NP_898870 7117 NM_004641 NP_904632 8028 NM_002823 NP_902814 5757 NM_002836 NP_902849 26212 NM_00636 NP_900627 648 NM_00636 NP_900827 644711 NM_000330 NP_9004072 1617 NM_001185 NP_900321 6247 NM_005266 NP_900527 2209 NM_0031483 NP_115684 84268 NM_004532 NP_901364 84268 NM_004532 NP_906520 2002 NM_033459 NP_976325 2710 NM_003459 NP_98234 23271 NM_005569 NP_98254 23271 NM_005569 NP_905560 3985 NM_000565 NP_900556 3570

synaptotagmin binding, cytoplasmic RNA	SYNCRIP	10492	NP_006363	NM_006372	4D	217834_s_at
NSFL1 (p97) cofactor (p47)	NSFLIC	55968	NP_057227	NM_016143	4D	217831_s_at
hypothetical protein FLJ13213	FLJ13213	79811	NP_079031	NM_024755	4D	217828_at
actin related protein 2/3 complex, subunit 4, 20kD	ARPC4	10093	NP_005709	NM_005718	4D	217818_s_at
suppressor of Ty 16 homolog (S. cerevisiae)	SUPT16H	11198	NP_009123	NM_007192	4D	217815_at
selenoprotein T	SELT	51714	NP_057359	NM_016275	4D	217811_at
Ufm1conjugating enzyme 1	Ufc1	51506	NP_057490	NM_016406	4D	217797_at
RABIIB, member RAS oncogene family	RAB11B	9230	NP_004209	NM_004218	4D	217793_at
G protein pathway suppressor 1	GPS1	2873	NP_004118	NM_004127	4D	217782_s_at
hypothetical protein HSPC152	HSPC152	51504	NP_057488	NM_016404	4D	217774_s_at
NADH dehydrogenase (ubiquinone) I alpha subcomplex	NDUFA4	4697	NP_002480	NM_002489	40	217773_s_at
chromosome 13 open reading frame 12	C13orf12	51371	NP_057016	NM_015932	4D	217769 s_at
chromosome 14 open reading frame 166	C14orf166	51637	NP_057123	NM_016039	4D	217768_at
small membrane protein 1	SMP1	23585	NP_055128	NM_014313	4D	217766_s_at
RAB31, member RAS oncogene family	RAB31	11031	NP_006859	NM_006868	4D	217764_s_at
RAB31, member RAS oncogene family	RAB31	11031	NP_006859	NM_006868	4D	217763_s_at
membranetype l matrix metalloproteinase cytoplasmi	MTCBPI	55256	NP_060739	NM_018269	45	217/61_at
tripartite motifcontaining 44	TRIM44	54765	NP_060053	NM_017583	4D	217759_at
small EDRKrich factor 2	SERF2	10169	NP_005761	NM_005770	4D	217756_x_at
hematological and neurological expressed 1	HNI	51155	NP_057269	NM_016185	4D	217755_at
adiponectin receptor 1	ADIPORI	51094	NP_057083	NM_015999	4D	217748_at
transmembrane protein 30A	TMEM30A	55754	NP_060717	NM_018247	4D	217743_s_at
preBcell colony enhancing factor 1	PBEF1	10135	NP_005737	NM_005746	4D	217739_s_at
preBcell colony enhancing factor 1	PBEF1	10135	NP_005737	NM_005746	4D	217738_at
WD repeat domain 6	WDR6	11180	NP_060501	NM_018031	4D	217734_s_at
thymosin, beta 10	TMSB10	9168	NP_066926	NM_021103	4D	217733_s_at
integral membrane protein 2B	ITM2B	9445	NP_068839	NM_021999	4D	217732_s_at
S100 calcium binding protein A6 (calcyclin)	S100A6	6277	NP_055439	NM_014624	4D	217728_at
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP_004198	NM_004207	4D	217691_x_at
winglesstype MMTV integration site family, member	WNT7B	7477	NP_478679	NM_058238	4D	217681_at
GTP binding protein 4	GTPBP4	23560	NP_036473	NM_012341	4D	217631_at
cation channel, sperm associated 2	CATSPER2	117155	NP_742095	NM_172097	4D	217588_at

	MBBI 13	20008	ND 05/1707	NM 014078	4	218049 s at
stannin	SNN	8303	NP_003489	NM_003498	4D	218033_s_at
stannin	SNN	8303	NP_003489	NM_003498	4D	218032_at
vaccinia related kinase 3	VRK3	51231	NP_057524	NM_016440	4D	1
chromosome 20 open reading frame 149	C20orf149	79144	NP_077275	NM_024299	4D	218010_x_at
ribosomal protein S27like	RPS27L	51065	NP_057004	NM_015920	4D	218007_s_at
zinc finger protein 22 (KOX 15)	ZNF22	7570	NP_008894	NM_006963	4D	218005_at
mitochondrial ribosomal protein S2	MRPS2	51116	NP_057118	NM_016034	4D	218001_at
sulfide quinone reductaselike (yeast)	SQRDL	58472	NP_067022	NM_021199	4D	217995_at
hypothetical protein FLJ20542	FLJ20542	54973	NP_060341	NM_017871	4D	217994 x at
cyclin B1 interacting protein 1	CCNB11P1	57820	NP_067001	NM_021178	4D	217988_at
selenoprotein X, 1	SEPX1	51734	NP_057416	NM_016332	4D	217977_at
mitogenactivated protein kinase kinase 1 interacti	MAP2K11P1	8649	NP_068805	NM_021970	4D	217971_at
nucleolar protein family A, member 3 (H/ACA small	NOLA3	55505	NP_061118	NM_018648	45	21/962_at
vitamin K epoxide reductase complex, subunit 1	VKORC1	79001	NP_076869	NM_024006	4D	217949_s_at
hypothetical protein FLJ10769	FLJ10769	55739	NP_060680	NM_018210	4D	217940 s_at
signal peptidase complex subunit 1 homolog (S	SPCS1	289/2	NF_034/60	NM_014041	45	21/92/_at
HSPC023 protein	HSPC023	28974	NP_054766	NM_014047	45	21/926_at
dynein, cytoplasmic, light polypeptide 2A	DNCL2A	83658	NP_054902	NM_014183	4D	217918_at
family with sequence similarity 49, member	FAM49B	51571	NP_057707	NM_016623	4D	217916_s_at
hect domain and RLD 2	HERC2	8924	NP_004658	NM_004667	4D	217902 s_at
chromosome 15 open reading frame 24	C15orf24	56851	NP_064539	NM_020154	4D	217898_at
NEFAinteracting nuclear protein NIP30	NIP30	80011	NP_079222	NM_024946	4D	217896_s_at
hypothetical protein FLJ13868	FLJ13868	64755	NP_073581	NM_022744	4D	217891_at
chromosome 2 open reading frame 25	C2orf25	27249	NP_056517	NM_015702	4D	217883_at
cell division cycle 27	CDC27	996	NP_001247	NM_001256	4D	217881_s_at
cell division cycle 27	CDC27	996	NP_001247	NM_001256	4D	217878_s_at
macrophage migration inhibitory factor (glycosylat	MIF	4282	NP_002406	NM_002415	4D	217871_s_at
Enah/Vasplike	EVL	51466	NP_057421	NM_016337	4D	217838_s_at
vacuolar protein sorting 24 (yeast)	VPS24	51652	NP_057163	NM_016079	4D	217837_s_at
chromosome 20 open reading frame 24	C20orf24	55969	NP_061328	NM_018840	4D	217835_x_at
Himacing					¥	

LanC lantibiotic synthetase component Clike 2	LANCL2	23913	INF_00110/	MINI_01909/	100	210217 3 at
Cili offiosoffie 11 open reading frame 10	CHOILIO	55015	MID 061167	NIM 010607		18210
Character 10	Cllorfin	746	ND 055021	NM 014206	4D	218213 s at
fructosamine3kinaserelated protein	FN3KRP	79672	NP_078895	NM_024619	4D	218210_at
nucleolar protein family 6 (RNAassociated)	NOL6	65083	NP_075068	NM_022917	4D	218199_s_at
osteopetrosis associated transmembrane protein 1	OSTMI	28962	NP_054747	NM_014028	4D	218196_at
ubiquinolcytochrome c reductase complex (7.2 kD)	HSPC051	29796	NP_037519	NM_013387	4D	218190_s_at
vacuolar protein sorting 4B (yeast)	VPS4B	9525	NP_004860	NM_004869	4D	218171_at
ADPribosylation factorlike 5	ARL5	26225	NP_036229	NM_012097	4D	218150_at
signal recognition particle receptor, B subunit	SRPRB	58477	NP_067026	NM_021203	4D	218140_x_at
NIF3 NGG1 interacting factor 3like 1 (S. pombe)	NIF3L1	60491	NP_068596	NM_021824	4D	218133_s_at
leukocyte receptor cluster (LRC) member 5	LENG5	79042	NP_076980	NM_024075	4D	218132_s_at
chromosome 21 open reading frame 59	C21orf59	56683	NP_060305	NM_017835	4D	218123_at
hom						
translocase of inner mitochondrial membrane 23	TIMM23	10431	NP 006318	NM_006327	4D	218118_s_at
ringbox 1	RBX1	9978	NP_055063	NM_014248	4D	218117_at
chromosome 9 open reading frame 78	C9orf78	51759	NP_057604	NM_016520	4D	218116_at
ASF1 antisilencing function 1 homolog B (S. cerevi	ASF1B	55723	NP_060624	NM_018154	45	218115_at
testis expressed sequence 10	TEX10	54881	NP_060216	NM_017746	40	218104_at
subcomplex unkn			***************************************			
NADH dehydrogenase (phiguinone) 1	NDUFC2	4718	NP 004540	NM 004549	4D	218101_s_at
uncharacterized hypothalamus protein HT008	HT008	55852	NP_060939	NM_018469	4D	218099_at
ADPribosylation factor guanine nucleotideexchange	ARFGEF2	10564	NP_006411	NM_006420	4D	218098_at
lacylglycerol3phosphate Oacyltransferase 5 (lysoph	AGPAT5	55326	NP_060831	NM_018361	40	218096_at
chromosome 20 open reading frame 35	C20orf35	55861	NP_060948	NM_018478	4D	218094_s_at
chromosome 20 open reading frame 4	C20orf4	25980	NP_056326	NM_015511	4D	218089_at
chromosome 20 open reading frame 27	C20orf27	54976	NP_060344	NM_017874	4D	218081_at
zinc finger, DHHC domain containing 3	ZDHHC3	51304	NP_057682	NM_016598	4D	218077 s_at
CG1128 protein	CG1128	51647	NP_057146	NM_016062	4D	218074_at
hypothetical protein FLJ10407	FLJ10407	55706	NP_060557	NM_018087	4D	218073_s_at
maleenhanced antigen	MEA	4201	NP_055438	NM_014623	4D	218061_at
HSPC038 protein	LOC51123	51123	NP_057180	NM_016096	4D	218059_at
bifunctional apoptosis regulator	BFAR	51283	NP_057645	NM_016561	4D	218056_at

translocase of inner mitochondrial membrane 8	TIMM8B	26521	NP_036591	NM_012459	4D	218357_s_at
FtsJ homolog 2 (E. coli)	FTSJ2	29960	NP_037525	NM_013393	4D	218356_at
Zwilch	FLJ10036	55055	NP_060445	NM_017975	4D	218349_s_at
retinoic acid induced 16	RAI16	64760	NP_073586	NM_022749	4D	218337_at
prefoldin 2	PFDN2	5202	NP_036526	NM_012394	4D	218336_at
brain expressed, Xlinked 1	BEXI	55859	NP_060946	NM_018476	4D	218332_at
coenzyme Q4 homolog (yeast)	COQ4	51117	NP_057119	NM_016035	4D	218328_at
spermatogenesis associated, serinerich 2	SPATS2	65244	NP_075559	NM_023071	4D	218324_s_at
pellino homolog 1 (Drosophila)	PELII	57162	NP_065702	NM_020651	4D	218319_at
translocase of inner mitochondrial membrane 9 homo	HMM9	20320	NF_030392	NW_012400	+	710010_at
CDK5 regulatory subunit associated protein 1	CDK5RAP1	51654	NP_057492	NM_016408	15	218315 s_at
calcium/calmodulindependent protein kinase II	CaMKIINalpha	55450	NP_061054	NM_018584	4D	at
				NM_018468	4D	218302_at
chromosome 10 open reading frame 97	C10orf97	80013	NP_079224	NM_024948	4D	218297_at
nucleoporin 50kDa	NUP50	10762	NP_009103	NM_007172	4D	218295_s_at
protein kinase, AMPactivated, gamma 2 noncatalytic	PRKAG2	51422	NP_05/28/	NM_016203	41)	218292_s_at
mitogenactivated proteinbinding proteininteracting	MAPBPIP	28956	NP_054736	NM_014017	4D	218291_at
ring finger protein 7	RNF7	9616	NP_055060	NM_014245	4D	218286_s_at
dehydrogenase/reductase (SDR family) member 6	DHRS6	56898	NP_064524	NM_020139	4D	218285_s_at
synovial sarcoma translocation gene on chromosome	SS18L2	51188	NP_057389	NM_016305	4D	218283_at
mitochondrial ribosomal protein L24	MRPL24	79590	NP_078816	NM_024540	4D	218270_at
BRCA2 and CDKN1A interacting protein	BCCIP	56647	NP_057651	NM_016567	4D	218264_at
transposonderived Buster1 transposaselike protein	LOC58486	58486	NP_067034	NM_021211	4D	218263_s_at
polymerase (RNA) I polypeptide D, 16kDa	POLRID	51082	NP_057056	NM_015972	4D	218258_at
fibrosin l	FBS1	64319	NP_071897	NM_022452	4D	218255_s_at
zinc finger, DHHC domain containing 6	ZDHHC6	64429	NP_071939	NM_022494	4D	218249_at
nucleolar protein 8	NOL8	55035	NP_060418	NM_017948	4D	218244_at
				NM_017601	4D	218233_s_at
pogo transposable element with KRAB domain	POGK	57645	NP_060012	NM_017542	4D	218229_s_at
NADH dehydrogenase (ubiquinone) 1 beta subcomplex,	NDUFB4	4710	NP_004538	NM_004547	4D	218226_s_at
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chromosome 21 open reading frame 66	C21orf66	94104	NP_057715	NM_016631	40	218515_at
hypothetical protein FLJ10587	FLJ10587	55181	NP_060619	NM_018149	4D	218514_at
WD repeat domain 12	WDR12	55759	NP_060726	NM_018256	4D	11
trichorhinophalangeal syndrome I	TRPSI	7227	NP_054831	NM_014112	4D	218502_s_at
ubiquitouslyexpressed transcript	UXT	8409	NP_004173	NM_004182	4D	218495_at
zinc finger protein 302	ZNF302	55900	NP_060913	NM_018443	4D	218490_s_at
hypothetical protein FLJ21827	FLJ21827	56912	NP_064538	NM_020153	4D	218483 s_at
exosome component 5	EXOSC5	56915	NP_064543	NM_020158	4D	218481_at
BardetBiedl syndrome 1	BBS1	582	NP_078925	NM_024649	4D	218471_s_at
CGI04 protein	CGI04	51067	NP_057020	NM_015936	4D	218470_at
tumor necrosis factor superfamily, member 5 induced	TNFSF5IP1	56984	NP_064617	NM_020232	4	21840/_80
MUS81 endonuclease homolog (yeast)	MUS81	80198	NP_079404	NM_025128	4D	218463_s_at
	MGC14560	51184	NP_057385	NM_016301	4D	218461_at
hypothetical protein FLJ20397	FLJ20397	54919	NP_060272	NM_017802	4D	218460_at
NFS1 nitrogen fixation 1 (S. cerevisiae)	NFS1	9054	NP_066923	NM_021100	4D	218455_at
hypothetical protein FLJ22662	FLJ22662	79887	NP_079105	NM_024829	4D	218454_at
CUB domaincontaining protein 1	CDCP1	64866	NP_073753	NM_022842	4D	218451_at
hypothetical protein FLJ11200	FLJ11200	55325	NP_060829	NM_018359	4D	218449_at
DC13 protein	DC13	56942	NP_064573	NM_020188	4D	218447_at
RNA polymerase II associated protein 1	RPAP1	26015	NP_056355	NM_015540	4D	218441_s_at
methylcrotonoylCoenzyme A carboxylase 1 (alpha)	MCCC1	56922	NP_064551	NM_020166	4D	218440_at
mediator of RNA polymerase II transcription, subun	MED28	80306	NP_079481	NM_025205	45	218458_s_at
serologically defined colon cancer antigen 3	SDCCAG3	10807	NP_006634	NM_006643	#D	218427_at
sorting nexin 10	SNX10	29887	NP_037454	NM_013322	4D	218404_at
zinc finger protein 281	ZNF281	23528	NP_036614	NM_012482	4D	218401_s_at
RNA binding motif protein 7	RBM7	10179	NP_057174	NM_016090	4D	218379_at
PRKR interacting protein 1 (IL11 inducible)	PRKRIP1	79706	NP_078929	NM_024653	4D	218378 s_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT9	53343	NP_076952	NM_024047	4D	218375_at
chromosome 12 open reading frame 4	C12orf4	57102	NP_065107	NM_020374	4D	218374_s_at
FLJ20859 gene	FLJ20859	64745	NP_073571	NM_022734	4D	218366 x_at
chromosome 14 open reading frame 114	C14orf114	55218	NP_060669	NM_018199	4D	218363_at

transducer of regulated cAMP response	TORC3	64784	NP_073606	NM_022769	4D	218648_at
hypothetical protein FLJ20534	FLJ20534	54969	NP_060337	NM_017867	4D	218646_at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	218643_s_at
coiledcoilhelixcoiledcoilhelix domain containing	CHCHD7	79145	NP_077276	NM_024300	4D	218642_s_at
				NM_023941	4D	218641_at
abhydrolase domain containing 10	ABHD10	55347	NP_060864	NM_018394	4D	218633_x_at
arginine vasopressininduced 1	AVPII	60370	NP_068378	NM_021732	4D	218631_at
smoothened homolog (Drosophila)	SMO	6608	NP_005622	NM_005631	4D	218629_at
hypothetical protein FLJ11259	FLJ11259	55332	NP_060840	NM_018370	4D	218627_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT2	318	NP_001152	NM_001161	4D	218609_s_at
SDA1 domain containing 1	SDADI	55153	NP_060585	NM_018115	4D	218607_s_at
transcription factor B2, mitochondrial	TFB2M	64216	NP_071761	NM_022366	4D	218605_at
LEM domain containing 3	LEMD3	23592	NP_055134	NM_014319	4D	218604_at
upregulated gene 4	URG4	55665	NP_060390	NM_017920	4D	218601_at
				NM_018072	4D	218594_at
progressive external ophthalmoplegia l	PEO1	56652	NP_068602	NM_021830	4D	218590_at
chromosome 5 open reading frame 3	C5orf3	10827	NP_061161	NM_018691	4D	218588_s_at
abhydrolase domain containing 4	ABHD4	63874	NP_071343	NM_022060	4D	218581_at
DEAH (AspGluAlaHis) box polypeptide 35	DHX35	60625	NP_068750	NM_021931	4D	218579_s_at
hyperparathyroidism 2 (with jaw tumor)	HRPT2	79577	NP_078805	NM_024529	4D	218578_at
anaphase promoting complex subunit 1	ANAPC1	64682	NP_073153	NM_022662	4D	218575_at
chromosome 14 open reading frame 123	C14orf123	29082	NP_054888	NM_014169	4D	218571_s_at
hypothetical protein FLJ10747	FLJ10747	55219	NP_060672	NM_018202	4D	218562_s_at
chromosome 6 open reading frame 149	C6orf149	57128	NP_065141	NM_020408	4D	218561_s_at
v-maf musculoaponeurotic fibrosarcoma oncogene homolog B (avian)	MAFB	9935	NP_005452	NM_005461	4D	218559_s_at
anaphase promoting complex subunit 2	ANAPC2	29882	NP_037498	NM_013366	4D	218555_at
enoyl Coenzyme A hydratase domain containing	ECHDC2	55268	NP_060751	NM_018281	4D	218552_at
MRS2like, magnesium homeostasis factor (S. cerevis	MRS2L	57380	NP_065713	NM_020662	4D	218538_s_at
hypothetical protein FLJ21749	FLJ21749	80194	NP_079400	NM_025124	4D	218531_at
aprataxin	APTX	54840	NP_060162	NM_017692	4D	218527_at
				NM_014185	4D	218526_s_at

FOSlike antigen 2	rOSL2	2333	TIME 000244	MINI OCCESS	-	
nyponiencai protein r 1312998	EOSI 2	2255	775500 div	NM 005253	B	3
hypothetical protein El 112000	FI 117998	64779	NP 073601	NM 022764	4D	218879 s at
hypothetical protein FLJ22390	FLJ22390	64757	NP_073583	NM_022746	4D	218865_at
WD repeat domain 33	WDR33	55339	NP_060853	NM_018383	4D	218851_s_at
hypothetical protein FLJ12788	FLJ12788	64427	NP_071937	NM_022492	4D	218838_s_at
phospholipid scramblase 3	PLSCR3	57048	NP_065093	NM_020360	4D	218828_at
aminopeptidaselike 1	NPEPL1	79716	NP_078939	NM_024663	4D	218822_s_at
				NM_025156	4D	218812_s_at
chromosome 7 open reading frame 19	C7orf19	80228	NP_116220	NM_032831	4D	218811_at
vav 3 oncogene	VAV3	10451	NP_006104	NM_006113	4D	218807_at
thioredoxinlike 4B	TXNL4B	54957	NP_060323	NM_017853	4D	218794_s_at
sex comb on midleglike 1 (Drosophila)	SCML1	6322	NP_006737	NM_006746	4D	218793_s_at
methionine sulfoxide reductase B2	MSRB2	22921	NP_036360	NM_012228	4D	218773_s_at
transmembrane protein 39B	ТМЕМЗ9В	55116	NP_060526	NM_018056	4D	218770_s_at
				NM_015940	4D	218760_at
hypothetical protein FLJ23323	FLJ23323	79707	NP_078930	NM_024654	4D	218754_at
hypothetical protein FLJ10307	FLJ10307	55113	NP_060523	NM_018053	4D	218753_at
hypothetical protein FLJ20422	FLJ20422	54929	NP_060284	NM_017814	4D	218745_x_at
chromosome 22 open reading frame 18	C22orf18	79019	NP_076958	NM_024053	4D	218741_at
ring finger protein 138	RNF138	51444	NP_057355	NM_016271	4D	218738_s_at
zinc finger protein 544	ZNF544	27300	NP_055295	NM_014480	4D	218735_s_at
latexin	LXN	56925	NP_064554	NM_020169	4D	218729 at
hypothetical protein FLJ12436	FLJ12436	79714	NP_078937	11	4D	218722_s_at
leprecanlike 1	LEPRELI	55214	NP_060662	NM_018192	4D	218717_s_at
hepatocellular carcinomaassociated antigen 66	HCA66	55813	NP_060898	NM_018428	4D	218715_at
NTF2like export factor 1	NXTI	29107	NP_037380	NM_013248	4D	218708_at
lactamase, beta 2	LACTB2	51110	NP_057111	NM_016027	4D	218701_at
RAB7, member RAS oncogene familylike 1	RAB7L1	8934	NP_003920	NM_003929	4D	218699_at
leucine rich repeat containing 5	LRRC5	55144	NP_060573	NM_018103	4D	218684_at
hypothetical protein FLJ13611	FLJ13611	80006	NP_079217	NM_024941	4D	218674_at
ATPase inhibitory factor 1	ATPIF1	93974	NP_057395	NM_016311	4D	218671 s_at
RAP2C, member of RAS oncogene family	RAP2C	57826	NP_067006	NM_021183	4D	218669_at
hypothetical protein FLJ14154	FLJ14154	79903	NP_079121	NM_024845	4D	218661_at
mitochondrial ribosomal protein S33	MRPS33	51650	NP_057155	NM_016071	4D	218654_s_at
				NM_022775	4D	218650_at

hypothetical protein FLJ14054	FLJ14054	79614	NP_078839	NM_024563	#5	219054_at
vacuolar protein sorting 37C (yeast)	VPS37C	55048	NP_060436	NM_017966	40	219053_s_at
coronin 7	CORO7	79585	NP_078811	NM_024535	4D	219040_at
CG1115 protein	CGI115	51018	NP_057136	NM_016052	4D	219037_at
poly (ADPribose) polymerase family, member 16	PARP16	54956	NP_060321	NM_017851	4D	219034_at
poly (ADPribose) polymerase family, member 8	PARP8	79668	NP_078891	NM_024615	4D	219033_at
opsin 3 (encephalopsin, panopsin)	OPN3	23596	NP_055137	NM_014322	4D	219032_x_at
CGI121 protein	CG1121	51002	NP_057142	NM_016058	4D	219030_at
myosin IXA	МҮО9А	4649	NP_008832	NM_006901	4D	219027_s_at
pleckstrin homology domain containing, family A (p	PLEKHAI	59338	NP_067635	NM_021622	4D	219024_at
nucleoporin 43kDa	NUP43	348995	NP_078923	NM_024647	4D	219007_at
chromosome 6 open reading frame 66	C6orf66	29078	NP_054884	NM_014165	4D	219006_at
hypothetical protein FLJ11000	FLJ11000	55281	NP_060765	NM_018295	4D	218999 at
RNA methyltransferase like 1	RNMTL1	55178	NP_060616	NM_018146	4D	218993_at
hypothetical protein FLJ20035	FLJ20035	55601	NP_060101	NM_017631	4D	218986_s_at
hypothetical protein FLJ20485	FLJ20485	54517	NP_061915	NM_019042	4D	218984_at
mitochondrial ribosomal protein S17	MRPS17	51373	NP_057053	NM_015969	4D	218982_s_at
ACN9 homolog (S. cerevisiae)	ACN9	57001	NP_064571	NM_020186	4D	218981_at
				NM_018586	4D	218978_s_at
zinc finger protein 64 homolog (mouse)	ZFP64	55734	NP_060667	NM_018197	4D	218968 s_at
hypothetical protein MGC3265	MGC3265	78991	NP_076933	NM_024028	4D	218953_s_at
glutaminyltRNA synthase (glutaminehydrolyzing)like	QRSL1	352/8	NP_060762	NIVI_018292	+	210747 3 41
phosphatidylinositol4phosphate 5kinase, type II, g	PIP5K2C	79837	NP_079055	NM_024779	1 =	218942_at
leucine zipperEFhand containing transmembrane prot	LETMI	3954	NP_036450	NM_012318	;	f 1
myoneurin	MYNN	55892	NP_061127	NM_018657	1	218926_at
chitobiase, diNacetyl	CTBS	1486	NP_004379	NM_004388	4D	218924 s_at
hypothetical protein FLJ14007	FLJ14007	79752	NP_078975	NM_024699	4D	218919_at
hypothetical protein FLJ10375	FLJ10375	55129	NP_060545	NM_018075	4D	218910_at
hypothetical protein FLJ20530	FLJ20530	55656	NP_060334	NM_017864	4D	218905_at
mitochondrial ribosomal protein 1.35	MRPL35	51318	NP_057706	NM_016622	4D	218890_x_at
MLF1 interacting protein	MLFIIP	79682	NP_078905	NM_024629	4D	218883_s_at
WD repeat domain 3	WDR3	10885	NP_006775	NM_006784	4D	218882_s_at

RAB8B, member RAS oncogene family	RAB8B	51762	NP_057614	NM_016530	[4D]	219210_s_at
interferon induced with helicase C domain 1	IFIHI	64135		NM_022168	4D	219209_at
ubiquitin associated protein 2	UBAP2	55833	NP_060919	NM_018449	4D	219192_at
eukaryotic translation initiation factor 2C, 4	EIF2C4	192670	NP_060099	NM_017629	4D	219190_s_at
pleckstrin homology, Sec7 and coiledcoil domains 4	PSCD4	27128	NP_037517	NM_013385	4D	219183_s_at
hypothetical protein FLJ22555	FLJ22555	79568	NP_078796	NM_024520	4D	219176_at
ubiquitin domain containing 1	UBTD1	80019	NP_079230	NM_024954	4D	219172_at
zinc finger protein 236	ZNF236	7776	NP_031371	NM_007345	4D	219171_s_at
mitochondrial ribosomal protein L11	MRPL11	65003	NP_057134	NM_016050	4D	219162 s at
kelchlike 2, Mayven (Drosophila)	KLHL2	111275	NP_009177	NM_007246	4D	219157_at
synaptojanin 2 binding protein	SYNJ2BP	55333	NP_060843	NM_018373	4D	219156_at
podocalyxinlike 2	PODXL2	50512	NP_056535	NM_015720	4D	219152_at
RAB, member of RAS oncogene familylike 2B	RABL2B	11158	NP_009012	NM_007081	4D	219151 s at
debranching enzyme homolog 1 (S. cerevisiae)	DBR1	51163	NP_057300	NM_016216	4D	219149_x_at
hypothetical protein FLJ22729	FLJ22729	79736	NP_078959	NM_024683	4D	219146_at
hypothetical protein FLJ20604	FLJ20604	54995	NP_060367	NM_017897	4D	219133_at
hypothetical protein FLJ10287	FLJ10287	54482	NP_061956	NM_019083	4D	219130_at
hypothetical protein FLJ20558	FLJ20558	54980	NP_060350	NM_017880	4D	219128_at
hypothetical protein MGC11242	MGC11242	79170	NP_077296	NM_024320	4D	219127_at
zinc finger protein 232	ZNF232	7775	NP_055334	NM_014519	4D	219123_at
hypothetical protein FLJ20171	FLJ20171	54845	NP_060167	NM_017697	4D	219121 s at
LSM8 homolog, U6 small nuclear RNA associated (S.	LSM8	51691	NP_057284	NM_016200	40	219119_at
DEAD (AspGluAlaAsp) box polypeptide 54	DDX54	79039	NP_076977	NM_024072	4D	219111_s_at
DEAD (AspGluAlaAsp) box polypeptide 27	DDX27	55661	NP_060365	NM_017895	4D	219108_x_at
	RNF141	50862	NP_057506	NM_016422	4D	219104_at
chromosome 12 open reading frame 5	C12orf5	57103	NP_065108	NM_020375	4D	219099_at
armadillo repeat containing 7	ARMC7	79637	NP_078861	NM_024585	4D	219096_at
hypothetical protein FLJ20701	FLJ20701	55022	NP_060403	NM_017933	4D	219093_at
NADPH cytochrome B5 oxidoreductase	NCB5OR	51167	NP_057314	NM_016230	4D	219079_at
fetal globininducing factor	FGIF	54851	NP_060174	NM_017704	4D	219069_at
chromosome 10 open reading frame 86	C10orf86	54780	NP_060085	NM_017615	4D	219067_s_at
hypothetical protein MDS018	MDS018	60490	NP_068595	NM_021823	4D	219066_at
hypothetical protein FLJ11712	FLJ11712	79621	NP_078846	NM_024570	4D	219056_at

endomucin	ENCN	21/02	141 027220	11111 010414		Sales and
uiggring receptor expressed on inveloid cells	EMON	50705	NID 057376	CPC910 WN	4D	219436 s at
triggering recentor expressed on mysloid calls 1	TREM!	54210	NP 061113	NM_018643	4D	219434_at
chromosome 18 open reading frame 22	C18orf22	79863	NP_079081	NM_024805	4D	219419_at
transmembrane protein 45A	TMEM45A	55076	NP_060474	NM_018004	4D	219410_at
cell deathinducing DFFAlike effector c	CIDEC	63924	NP_071377	NM_022094	4D	219398_at
hypothetical protein FLJ13448	FLJ13448	80219	NP_079423	NM_025147	4D	219397_at
NMDA receptor regulated 1 like	NARGIL	79612	NP_078837	NM_024561	4D	219378_at
chromosome 18 open reading frame 11	C18orf11	64762	NP_073588	NM_022751	4D	219377_at
dolichylphosphate mannosyltransferase polypeptide	DPM3	54344	NP_061846	NM_018973	45	2195/5_at
neuropilin 2	NRP2	8828	NP_061004	NM_018534	4D	219367_s_at
CGI12 protein	CGH2	51001	NP_057026	NM_015942	4D	219363_s_at
SNF7 domain containing 2	SNF7DC2	51510	NP_057494	NM_016410	4D	219356_s_at
CGI143 protein	CGI143	51027	NP_057158	NM_016074	4D	219345_at
activating signal cointegrator I complex subunit	ASCC1	51008	NP_057031	NM_015947	4D	219336_s_at
				NM_024062	4D	219330_at
chromosome 2 open reading frame 28	C2orf28	51374	NP_057169	NM_016085	4D	219329_s_at
mediator of RNA polymerase II transcription, subun	MED31	51003	NP_057144	NM_016060	40	219518_x_at
zinc finger and BTB domain containing 10	ZBTB10	65986	NP_076418	NM_023929	4D	219312_s_at
Fbox protein 2	FBXO2	26232	NP_036300	NM_012168	4D	219305_x_at
THAP domain containing, apoptosis associated prote	THAP1	55145	NP_060575	NM_018105	4D	219292_at
				NM_017718	4D	219279_at
chromosome 9 open reading frame 82	C9orf82	79886	NP_079104	NM_024828	4D	219276_x_at
hypothetical protein FLJ21616	FLJ21616	79618	NP_078843	NM_024567	4D	219269_at
zinc finger protein 350	ZNF350	59348	NP_067645	NM_021632	4D	219266_at
THUMP domain containing 2	THUMPD2	80745	NP_079540	NM_025264	4D	219248_at
mitochondrial ribosomal protein L46	MRPL46	26589	NP_071446	NM_022163	4D	219244_s_at
GTPase, IMAP family member 4	GIMAP4	55303	NP_060796	NM_018326	4D	219243_at
centrosome protein Cep63	Cep63	80254	NP_079456	NM_025180	4D	219242_at
secernin 3	SCRN3	79634	NP_078859	NM_024583	4D	219234_x_at
gasderminlike	GSDML	55876	NP_061000	NM_018530	4D	219233_s_at
zinc finger protein 331	ZNF331	55422	NP_061025	NM_018555	4D	219228_at
mitochondrial ribosomal protein S22	MRPS22	56945	NP_064576	NM_020191	4D	219220_x_at

Kruppellike factor 3 (basic)	KLF3	51274	NP_057615	NM_016531	4D	219657_s_at
protocadherin 12	PCDH12	51294	NP_057664	NM_016580	4D	219656_at
				NM_014054	4D	219653_at
developmental pluripotency associated 4	DPPA4	55211	NP_060659	NM_018189	4D	219651_at
popeye domain containing 2	POPDC2	64091	NP_071418	NM_022135	4D	219647_at
NYREN58 antigen	NYREN58	51134	NP_057206	NM_016122	4D	219644_at
deetiolated homolog 1 (Arabidopsis)	DET1	55070	NP_060466	NM_017996	4D	219641_at
				NM_024937	4D	219631_at
sirtuin (silent mating type information regulation	SIRT6	51548	NP_057623	NM_016539	4D	219613_s_at
hypothetical protein DKFZp434L0117	DKFZP434L011	64793	NP_073615	NM_022778	4D	219611_s_at
membranespanning 4domains, subfamily A, member 4	MS4A4A	51338	NP_076926	NM_024021	4D	219607_s_at
RWD domain containing 1	RWDD1	51389	NP_057188	NM_016104	4D	219598_s_at
ninjurin 2	NINJ2	4815	NP_057617	NM_016533	4D	219594_at
CGI30 protein	CGI30	51611	NP_057042	NM_015958	4D	219590_x_at
likely homolog of yeast SEN2	SEN2L	80746	NP_079541	NM_025265	4D	219581_at
reticulon 3	RTN3	10313	NP_006045	NM_006054	4D	219549_s_at
FLJ22624 protein	FLJ22624	79866	NP_079084	NM_024808	4D	219544_at
zinc finger protein 267	ZNF267	10308	NP_003405	NM_003414	4D	219540_at
chromosome 14 open reading frame 169	C14orf169	79697	NP_078920	NM_024644	4D	219526_at
transient receptor potential cation channel, subfa	TRPV4	59341	NP_067638	NM_021625	4D	219516_at
hypothetical protein FLJ13150	FLJ13150	79871	NP_079089	NM_024813	4D	219504_s_at
transmembrane protein 40	TMEM40	55287	NP_060776	NM_018306	4D	219503_s_at
Bcell CLL/lymphoma 11A (zinc finger protein)	BCL11A	53335	NP_060484	NM_018014	4D	219498_s_at
cysteinerich hydrophobic domain 2	CHIC2	26511	NP_036242	NM_012110	4D	219492_at
hypothetical protein FLJ23560	FLJ23560	79738	NP_078961	NM_024685	4D	219487_at
chromosome 21 open reading frame 18	C21orf18	54093	NP_059134	NM_017438	4D	219482_at
<u>I</u>				NM_024115	4D	219476_at
ganglioside induced differentiation associated pro	GDAP2	54834	NP_060156	NM_017686	4D	219473_at
hypothetical protein FLJ20125	FLJ20125	54826	NP_060146	NM_017676	4D	219467_at
polymerase (RNA) III (DNA directed) polypeptide	POLR3B	55703	NP_060552	NM_018082	4D	219459_at
				NM_016064	4D	219451_at
hypothetical protein MGC3020	MGC3020	79014	NP_076953	NM_024048	4D	219442_at

FYVE, RhoGEF and PH domain containing 6	FGD6	55785	NP_060821	NM_018351	40	219901_at
Ctype lectin domain family 5, member A	CLEC5A	23601	NP_037384	NM_013252	45	219890_at
hypothetical protein MGC3130	MGC3130	78995	NP_076937	NM_024032	4D	219879_s_at
hypothetical protein FLJ13842	FLJ13842	79698	NP_078921	NM_024645	4D	219877_at
activating transcription factor 7 interacting prot	ATF7IP2	80063	NP_079273	NM_024997	4D	219870_at
nuclear prelamin A recognition factor	NARF	26502	NP_036468	NM_012336	4D	219862_s_at
Ctype lectin domain family 4, member E	CLEC4E	26253	NP_055173	NM_014358	4D	219859_at
zinc finger protein 14 (KOX 6)	ZNF14	7561	NP_066358	NM_021030	4D	219854_at
hypothetical protein FLJ23506	FLJ23506	79891	NP_079109	NM_024833	4D	219849_at
				NM_025174	4D	219846_at
mitochondrial translational release factor 1	MTRF1	9617	NP_004285	NM_004294	4D	219822_at
mitochondrial ribosomal protein S28	MRPS28	28957	NP_054737	NM_014018	4D	219819_s_at
G patch domain containing 1	GPATC1	55094	NP_060495	NM_018025	4D	219818_s_at
apoptosisrelated protein PNAS1	FLJ39616	51275	NP_057618	NM_016534	4D	219817_at
complexinterac						
valosincontaining protein (p97)/p47	VCIP135	80124	NP_079330	NM_025054	4D	219810_at
FN5 protein	FN5	56935	NP_064564	NM_020179	4D	219806_s_at
paired immunoglobinlike type 2 receptor alpha	PILRA	29992	NP_038467	NM_013439	4D	219788_at
epithelial cell transforming sequence 2 oncogene	ECT2	1894	NP_060568	NM_018098	4D	219787_s_at
hypothetical protein FLJ10996	FLJ10996	54520	NP_061917	NM_019044	4D	219774_at
glycosyltransferaselike domain containing 1	GTDC1	79712	NP_078935	NM_024659	4D	219770_at
immune costimulatory protein B7H4	B7H4	79679	NP_078902	NM_024626	4D	219768_at
crystallin, zeta (quinone reductase)like 1	CRYZLI	9946	NP_005102	NM_005111	4D	219767_s_at
hypothetical protein MGC4093	MGC4093	80776	NP_085055	NM_030578	4D	219766_at
hypothetical protein FLJ12586	FLJ12586	79673	NP_078896	NM_024620	4D	219765_at
ribosomal protein L36	RPL36	25873	NP_056229	NM_015414	4D	219762_s_at
hypothetical protein FLJ21148	FLJ21148	79918	NP_079136	NM_024860	4D	219751_at
zinc finger protein 552	ZNF552	79818	NP_079038	NM_024762	4D	219741_x_at
protocadherin 9	PCDH9	5101	NP_065136	NM_020403	4D	219737_s_at
SID1 transmembrane family, member 1	SIDT1	54847	NP_060169	NM_017699	4D	219734_at
				NM_014796	4D	219724_s_at
hypothetical protein FLJ20280	FLJ20280	54876	NP_060211	NM_017741	4D	219717_at
plexin domain containing 1	PLXDC1	57125	NP_065138	NM_020405	4D	219700_at
hypothetical protein FLJ22573	FLJ22573	79713	NP_078936	NM_024660	4D	219690_at
RAB11 family interacting protein 1 (class I)	RAB11FIP1	80223	NP_079427	NM_025151	4D	219681_s_at

NP_079457 80255 SLC35F5 NP_004812 9421 HANDI NP_037455 29916 SNX11
55576 80255
060208 54875 C9orf39
060055 11182 SLC2A6
NP_001727 728 C5R1
29802 VPREB3
51438 MAGEC2
26228 BRDG1
057668 51561 IL23A
26277 TINF2
23409 SIRT4
23678 SGKL
55716 LIMR
23225 NUP210
56605 ERO1LB
23569 PADI4
8778 SIGLEC5
29094 HSPC159
64708 COPS7B
50615 IL21R
59272 ACE2
56672 C11orf17
50856 CLEC4A
51022 GLRX2
4054 LTBP3

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solute carrier family 30 (zinc transporter), membe	SLC30A10	55532	NP_061183	NM_018713	4D	220435_at
hypothetical protein MGC5356	MGC5356	79025	NP_076964	NM_024059	4D	220426_at
ubiquitin specific protease 25	USP25	29761	NP_037528	NM_013396	4D	220419 s_at
ubiquitin associated and SH3 domain containing,	UBASH3A	53347	NP_061834	NM_018961	4D	220418_at
				NM_014076	4D	220404_at
Mdm4, transformed 3T3 cell double minute 1, p53 bi	MDMI	56890	NP_064513	NM_020128	45	220397_at
zinc finger and BTB domain containing 3	ZBTB3	79842	NP_079060	NM_024784	4D	220391_at
leucine rich repeat containing 19	LRRC19	64922	NP_075052	NM_022901	4D	220376_at
mSin3Aassociated protein 130	SAP130	79595	NP_078821	NM_024545	4D	220367_s_at
polybromo l	PB1	55193	NP_060635	NM_018165	4D	220355_s_at
leucine rich repeat transmembrane neuronal 4	LRRTM4	80059	NP_079269	NM_024993	4D	220345_at
truncated calcium binding protein	LOC51149	51149	NP_057259	NM_016175	4D	220341_s_at
SAM domain, SH3 domain and nuclear localisation si	SAMSNI	64092	NP_0/1419	NM_022136		22030_S_at
chromosome 6 open reading frame 96	C6orf96	55005	NP_060379	NM_017909	1 1 1 1 1	220329_s_at
colon carcinoma related protein	FLJ38507	389136	NP_057290	NM_016206	4D	220327_at
neuronal PAS domain protein 3	NPAS3	64067	NP_071406	NM_022123	4D	220316_at
hypothetical protein FLJ13840	FLJ13840	79802	NP_079022	NM_024746	4D	220283_at
RIC3 protein	RIC3	79608	NP_078833	NM_024557	4D	220282_at
zinc finger, DHHC domain containing 4	ZDHHC4	55146	NP_060576	NM_018106	4D	220261_s_at
hypothetical protein FLJ10891	FLJ10891	55762	NP_060730	NM_018260	4D	220242_x_at
chromosome 13 open reading frame 11	C13orf11	55002	NP_060375	NM_017905	4D	220241_at
chromosome 13 open reading frame 11	C13orf11	55002	NP_060375	NM_017905	4D	220240_s_at
receptorinteracting factor 1	RIF1	55791	NP_060842	NM_018372	4D	220235_s_at
				NM_018692	4D	220213_at
thyroid adenoma associated	THADA	63892	NP_071348	NM_022065	4D	220212_s_at
zinc finger, MYM domain containing 1	ZMYMI	79830	NP_079048	NM_024772	4D	220206_at
methylCpG binding domain protein 5	MBD5	55777	NP_060798	NM_018328	4D	220195_at
nucleotide binding proteinlike	NUBPL	80224	NP_079428	NM_025152	4D	220176_at
				NM_020667	4D	220175_s_at
hypothetical protein FLJ11767	FLJ11767	79645	NP_078869	NM_024593	4D	220156_at
hypothetical protein FLJ22671	FLJ22671	79919	NP_079137	NM_024861	4D	220149_at
aldehyde dehydrogenase 8 family, member A1	ALDH8A1	64577	NP_072090	NM_022568	40	220148_at

solute carrier family 9 (sodium/hydrogen	SLC9A5	6553	NP_004585	NM_004594	40	220600_S_at
leucine rich repeat containing 31	LRRC31	79782	NP_079003	NM_024727	E	220622_at
forkhead box E3	FOXE3	2301		NM_012186	4D	220621_at
chromosome 1 open reading frame 42	C1orf42	54544	NP_061933	NM_019060	4D	220620_at
zinc finger protein 532	ZNF532	55205	NP_060651	NM_018181	4D	220617_s_at
male sterility domain containing 1	MLSTD1	55711	NP_060569	NM_018099	4D	220615_s_at
leucine rich repeat (in FLII) interacting protein	LRRFIP2	9209	NP_006300	NM_006309	4D	220610_s_at
ADPribosylationlike factor 6 interacting protein 4	ARL6IP4	51329	NP_061164	NM_018694	4D	220597_s_at
uncharacterized hematopoietic stem/progenitor cell	MDS028	55846	NP_060933	NM_018463	4D	220590_at
				NM_025086	4D	220583_at
				NM_025071	4D	220582_at
resistin	RETN	56729	NP_065148	NM_020415	4D	220570_at
phosphoinositide3kinase, regulatory subunit 5, p10	PIK3R5	23533	NP_055123	NM_014308	4D	220566_at
panhematopoietic expression	PHEMX	10077	NP_005696	NM_005705	4D	220558_x_at
family with sequence similarity 35, member A	FAM35A	54537	NP_061927	NM_019054	4D	220547_s_at
chromosome 10 open reading frame 92	C10orf92	54777	NP_060079	NM_017609	4D	220539_at
myotubularin related protein 8	MTMR8	55613	NP_060147	NM_017677	4D	220537_at
				NM_024853	4D	220533_at
mitochondrial ribosomal protein L20	MRPL20	55052	NP_060441	NM_017971	4D	220526_s_at
deleted in liver cancer 1	DLC1	10395	NP_079043	NM_024767	4D	220512_at
Teomplex protein 1	CESK1	150160	NP_055221	NM_014406	4D	220508_at
hepcidin antimicrobial peptide	HAMP	57817	NP_066998	NM_021175	4D	220491_at
				NM_018565	4D	220490_at
				NM_017698	4D	220486_x_at
deafness locus associated putative guanine nucleot	DELGEF	26297	NP_036271	NM_012139	4D	220482_s_at
G proteincoupled receptor 75	GPR75	10936	NP_006785	NM_006794	4D	220481_at
hypothetical protein LOC55924	LOC55924	55924	NP_061972	NM_019099	4D	220476_s_at
myc target 1	MYCT1	80177	NP_079383	NM_025107	4D	220471_s_at
				NM_025032	4D	220467_at
solute carrier organic anion transporter family, m	SLC01C1	53919	NP_059131	NM_017435	4D	220460_at
histamine receptor H3	HRH3	11255	NP_009163	NM_007232	4D	220447_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST4	10164	NP_005760	NM_005769	4D	220446_s_at
ventral anterior homeobox 2	VAX2	25806	NP_036608	NM_012476	4D	220443_s_at
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tigger transposable element derived 6	TIGD6	81789	NP_112215	NM_030953	40	220986_s_at
ring finger protein 170	RNF170	81790	NP_112216	NM_030954	4D	220985_s_at
sprouty homolog 4 (Drosophila)	SPRY4	81848	NP_112226	NM_030964	4D	220983_s_at
erythrocyte membrane protein band 4.1 like 5	EPB41L5	57669	NP_065960	NM_020909	4D	220977_x_at
keratin associated protein 11	KRTAP11	81851	NP_112229	NM_030967	4D	220976_s_at
interleukin 17E	IL17E	64806	NP_073626	NM_022789	4D	220971_at
				NM_030892	4D	220969_s_at
ribosomal protein L22	RPL22	6146	NP_000974	NM_000983	4D	220960_x_at
MANSC domain containing 1	MANSC1	54682	NP_060520	NM_018050	4D	220945_x_at
				NM_025190	4D	220940_at
				NM_024084	4D	220934_s_at
heparanase 2	HPSE2	60495	NP_068600	NM_021828	4D	220927_s_at
G proteincoupled receptor 157	GPR157	80045	NP_079256	NM_024980	4D	220901_at
PR domain containing 12	PRDM12	59335	NP_067632	NM_021619	4D	220894_x_at
DEAD (AspGluAlaAsp) box polypeptide 47	DDX47	51202	NP_057439	NM_016355	4D	220890_s_at
				NM_018614	4D	220883_at
				NM_018575	4D	220874_at
0				NM_014123	4D	220854_at
FERM domain containing 1	FRMDI	79981	NP_079195	NM_024919	4D	220819_at
gremlin 2 homolog, cysteine knot superfamily (Xeno	GREM2	64388	NP_071914	NM_022469	4D	220794_at
urotensin 2	UTS2	10911	NP_068835	NM_021995	4D	220785_at
urotensin 2	UTS2	10911	NP_068835	NM_021995	4D	220784 s_at
potassium inwardlyrectifying channel, subfamily J,	KCNJ14	3770	NP_037480	NM_013348	4D	220776_at
dymeclin	DYM	54808	NP_060123	NM_017653	4D	220774_at
chromosome 6 open reading frame 48	C6orf48	50854	NP_058643	NM_016947	4D	220755_s_at
zinc finger protein 580	ZNF580	51157	NP_057286	NM_016202	4D	220748_s_at
WD repeat domain 10	WDR10	55764	NP_060732	NM_018262	4D	220744_s_at
inorganic pyrophosphatase 2	PPA2	27068	NP_008834	NM_006903	4D	220741_s_at
				NM_024992	4D	220715_at
				NM_018563	4D	220704_at
				NM_018616	4D	220702_at
KIAA1622	KIAA1622	57718	NP_066009	NM_020958	4D	220673_s_at
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cyclin L2	CCNL2	81669	NP_112199	NM_030937	4D	221427_s_at
calcium channel, voltagedependent, gamma subunit 5	CACNG5	27091	NP_055219	NM_014404	4D	221401_at
ectodysplasin A2 receptor	EDA2R	60401	NP_068555	NM_021783	4D	221399_at
				NM_020527	4D	221379_at
recombining binding protein suppressor of hairless	RBPSUHL	11317	NP_055091	NM_014276	4D	221377_s_at
motilin receptor	MLNR	2862	NP_001498	NM_001507	4D	221365_at
hypothetical protein dJ122O8.2	DJ12208.2	57226	NP_065199	NM_020466	4D	221311_x_at
fibroblast growth factor receptor substrate 2	FRS2	10818	NP_006645	NM_006654	4D	221308_at
G proteincoupled receptor 22	GPR22	2845	NP_005286	NM_005295	4D	221288_at
pseudouridylate synthase 3	PUS3	83480	NP_112597	NM_031307	4D	221277_s_at
				NM_030825	4D	221235_s_at
hypothetical protein FLJ20628	FLJ20628	55006	NP_060380	NM_017910	4D	221229_s_at
hypothetical protein FLJ23342	FLJ23342	79684	NP_078907	NM_024631	4D	221208_s_at
YEATS domain containing 2	YEATS2	55689	NP_060493	NM_018023	4D	221203_s_at
chromosome 18 open reading frame 8	C18orf8	29919	NP_037458	NM_013326	4D	221190_s_at
				NM_025050	4D	221179_at
				NM_025043	4D	221177_at
histamine receptor H4	HRH4	59340	NP_067637	NM_021624	4D	221169_s_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST5	23563	NP_036258	NM_012126	4D	221164_x_at
achaetescute complex (Drosophila) homologlike 3	ASCL3	56676	NP_065697	NM_020646	#D	221161_at
cell cycle progression 1	CCPG1	9236	NP_004739	NM_004748	4D	221156_x_at
HT001 protein	HT001	28990	NP_054784	NM_014065	4D	221135_s_at
angiopoietin 4	ANGPT4	51378	NP_057069	NM_015985	4D	221134_at
hypothetical protein FLJ10826	FLJ10826	55239	NP_060703	NM_018233	4D	221090_s_at
family with sequence similarity 31, member C	FAM31C	79958	NP_079174	NM_024898	4D	221080_s_at
armadillo repeat containing 4	ARMC4	55130	NP_060546	NM_018076	4D	221077_at
spermatogenesis associated 1	SPATAI	64173	NP_071749	NM_022354	4D	221057_at
anterior pharynx defective 1 homolog B (C. elegans	APHIB	83464	NP_112591	NM_031301	4D	221036_s_at
transmembrane protease, serine 5 (spinesin)	TMPRSS5	80975	NP_110397	NM_030770	4D	221032_s_at
RAB33B, member RAS oncogene family	RAB33B	83452	NP_112586	NM_031296	4D	221014_s_at
chromosome I open reading frame 14	Clorf14	81626	NP_112195	NM_030933	4D	220996_s_at
likely ortholog of rat vacuole membrane protein 1	VMP1	81671	NP_112200	NM_030938	4D	220990_s_at

abhydrolase domain containing 6	ABHD6	57406	NP_065727	NM_020676	4D	221679_s_at
coronin, actin binding protein, 1C	COROIC	23603	NP_055140	NM_014325	4D	221676_s_at
LIM homeobox 3	LHX3	8022	NP_055379	NM_014564	4D	221670_s_at
likely ortholog of mouse synembryn	RIC8	60626	NP_068751	NM_021932	4D	221647_s_at
arginineglutamic acid dipeptide (RE) repeats	RERE	473	NP_036234	NM_012102	4D	221643_s_at
syntaxin 16	STX16	8675	NP_001001433	NM_001001433	4D	221638_s_at
uncharacterized hypothalamus protein HT007	HT007	55863	NP_060950	NM_018480	4D	221622_s_at
signaltransducing adaptor protein2	STAP2	55620	NP_060190	NM_017720	4D	221610_s_at
HSPC171 protein	HSPC171	29100	NP_054906	NM_014187	4D	221597_s_at
histone 3, H2a	HIST3H2A	92815	NP_254280	NM_033445	4D	221582_at
sterol Oacyltransferase (acylCoenzyme A: cholester	SOATI	6646	NP_003092	NM_003101	4D	221561_at
lymphoid enhancerbinding factor 1	LEF1	51176	NP_057353	NM_016269	4D	221558_s_at
cysteinerich secretory protein LCCL domain contain	CRISPLD2	83716	NP_113664	NM_031476	4D	221541_at
general transcription factor IIH, polypeptide 2, 4	GTF2H2	2966	NP_001506	NM_001515	4D	221540_x_at
hypothetical protein FLJ11301	FLJ11301	55341	NP_060855	NM_018385	4D	221536_s_at
ankyrin repeat domain 27 (VPS9 domain)	ANKRD27	84079	NP_115515	NM_032139	4D	221522_at
cell division cycle associated 8	CDCA8	55143	NP_060571	NM_018101	4D	221520_s_at
cofactor required for Sp1 transcriptional activati	CRSP6	9440	NP_004259	NM_004268	4D	221517_s_at
hypothetical protein FLJ20232	FLJ20232	54471	NP_061881	NM_019008	4D	221516_s_at
cell cycle progression 1	CCPG1	9236	NP_065790	NM_020739	4D	221511_x_at
karyopherin alpha 3 (importin alpha 4)	KPNA3	3839	NP_002258	NM_002267	4D	221503_s_at
eukaryotic translation initiation factor 3, subuni	EIF3S12	27335	NP_037366	NM_013234	4D	221494_x_at
sprouty homolog 4 (Drosophila)	SPRY4	81848	NP_112226	NM_030964	4D	221489_s_at
UDPGal:betaGlcNAc beta 1,4 galactosyltransferase,	B4GALT5	9334	NP_004/6/	NM_004//6	40	221485_at
galactosyltransferase,			——————————————————————————————————————	-		· ·
UDPGal:hetaGlcNAc heta 1.4	B4GALT5	9334	NP 004767	NM 004776	4D	221484 at
BCL2/adenovirus E1B 19kDa interacting protein 31ik	BNIP3L	060	NP_004322	NM_004331	4	2214/8_at
glucose6phosphatase, catalytic, 2	G6PC2	57818	NP_066999	NM_021176	4D	221453_at
olfactory receptor, family 2, subfamily W, member	OR2W1	26692	NP_112165	NM_030903	4D	221451_s_at
Tcell immunomodulatory protein	CDA08	81533	NP_110417	NM_030790	4D	221449_s_at
chromosome 14 open reading frame 156	C14orf156	81892	NP_112487	NM_031210	4D	221434_s_at

dom dom	LRCH4	4034	NF_002310	KI _002319	40	222011_X_at
tcomplex 1	TCP1	6950	NP_110379	NM_030752	3 6	222010_at
hypothetical protein FLJ10534	FLJ10534	55720		NM_018128	4D	221987_s_at
centaurin, gammalike family, member 1	CTGLFI	119016	NP_597703	NM_133446	4D	221971_x_at
ubiquitinconjugating enzyme E2H (UBC8 homolog, yea	UBE2H	7328	NP_003335	NM_003344	4D	221962_s_at
hypothetical protein LOC283232	LOC283232	283232	NP_777600	NM_174940	4D	221951_at
chromosome 14 open reading frame 87	C14orf87	51218	NP_057501	NM_016417	4D	221932_s_at
	DKFZp76112123	83637	NP_777589	NM_174929	4D	221924_at
hypothetical protein MGC1203	MGC1203	79140	NP_077272	NM_024296	4D	221912_s_at
likely ortholog of mouse hypoxia induced gene 1	HIGI	25994	NP_054775	NM_014056	4D	221896_s_at
ecotropic viral integration site 1	EVII	2122	NP_005232	NM_005241	4D	221884_at
major histocompatibility complex, class I, F	HLAF	3134	NP_061823	NM_018950	4D	221875_x_at
TBC1 domain family, member 12	TBCID12	23232	XP_051081	XM_051081	4D	221858_at
plakophilin I (ectodermal dysplasia/skin fragility	PKP1	5317	NP_001005337	NM_001005337	4D	221854_at
centaurin, gammalike family, member 1	CTGLF1	119016	NP_597703	NM_133446	4D	221850_x_at
RAP2A, member of RAS oncogene family	RAP2A	5911	NP_066361	NM_021033	4D	221830_at
hypothetical protein FLJ20436	FLJ20436	54934	NP_060292	NM_017822	4D	221821_s_at
KIAA1698 protein	KIAA1698	80789	NP_085131	NM_030628	4D	221818_at
PHD finger protein 11	PHF11	51131	NP_057203	NM_016119	4D	221816_s_at
RAB9A, member RAS oncogene family	RAB9A	9367	NP_004242	NM_004251	4D	221808_at
nuclear receptor binding factor 2	NRBF2	29982	NP_110386	NM_030759	4D	221803_s_at
ribosomal protein L22	RPL22	6146	NP_000974	NM_000983	4D	221775_x_at
jumonji domain containing 1C	JMJDIC	221037	NP_004232	NM_004241	4D	221763_at
YTH domain family, member 1	YTHDF1	54915	NP_060268	NM_017798	4D	221741_s_at
collagen, type V, alpha 2	COL5A2	1290	NP_000384	NM_000393	4D	221729_at
Ctype lectin domain family 4, member A	CLEC4A	50856	NP_057268	NM_016184	4D	221724_s_at
solute carrier family 4, sodium bicarbonate cotran	SLC4A5	57835	NP_067019	NM_021196	4D	221723_s_at
hypothetical protein FLJ10439	FLJ10439	54663	NP_060563	NM_018093	4D	221712_s_at
hypothetical protein FLJ12750	FLJ12750	79720	NP_078943	NM_024667	4D	221704_s_at
BBPlike protein 2	BLP2	80213	NP_510883	NM_078474	4D	221702_s_at
ubiquitin A52 residue ribosomal protein fusion pro	UBA52	7311	NP_003324	NM_003333	4D	221700_s_at
mitochondrial ribosomal protein L34	MRPL34	64981	NP_076426	NM_023937	4D	221692_s_at
Down syndrome critical region gene 5	DSCR5	51227	NP_710149	NM_153682	4D	221689_s_at

ankyrin repeat and SOCS boxcontaining 6	ASB6	140459	NP_060343	NM_017873	4D	222524_s_at
adaptorrelated protein complex 3, mu 1 subunit	AP3M1	26985	NP_996895	NM_207012	4D	222516_at
NEDD8 ultimate buster1	NYREN18	51667	NP_057202	NM_016118	4D	222512_at
chromosome 7 open reading frame 2	C7orf2	64327	NP_071903	NM_022458	4D	222505_at
testis expressed sequence 27	TEX27	60685	NP_068762	NM_021943	4D	222493_s_at
ubiquitinconjugating enzyme E2Q (putative)	UBE2Q	55585	NP_060052	NM_017582	4D	222480_at
translocase of outer mitochondrial membrane 22 hom	TOMM22	56993	NP_064628	NM_020243	4D	222474_s_at
subun						CHARGE STATE OF THE STATE OF TH
mediator of RNA polymerase II transcription	MED4	29079	NP 054885	NM 014166	4D	a l
vacuolar protein sorting 24 (yeast)	VPS24	51652	NP 057163	NM 016079	4D	222436 s at
coronin, actin binding protein, 1C	COROIC	23603	NP_055140	NM_014325	4D	222409_at
yippeelike 5 (Drosophila)	YPEL5	51646	NP_057145	190910_MN	4D	222408_s_at
butyrateinduced transcript 1	HSPC121	51495	NP_057479	NM_016395	4D	222404_x_at
mitochondrial carrier homolog 2 (C. elegans)	MTCH2	23788	NP_055157	NM_014342	4D	222403_at
small membrane protein 1	SMP1	23585	NP_055128	NM_014313	4D	222401_s_at
hematological and neurological expressed 1	HNI	51155	NP_057269	NM_016185	4D	222396_at
programmed cell death 6 interacting protein	PDCD6IP	10015	NP_037506	NM_013374	4D	222394_at
CGI30 protein	CG130	51611	NP_057042	NM_015958	4D	222360_at
splicing factor, arginine/serinerich 15	SFRS15	57466	NP_065757	NM_020706	4D	222311_s_at
sirtuin (silent mating type information regulation	SIRT4	23409	NP_036372	NM_012240	4D	222248_s_at
paired immunoglobinlike type 2 receptor alpha	PILRA	29992	NP_840057	NM_178273	4D	222218_s_at
LAG1 longevity assurance homolog 2 (S. cerevisiae)	LASS2	29956	NP_037516	NM_013384	4D	222212_s_at
hypothetical protein FLJ22104	FLJ22104	65084	NP_075069	NM_022918	4D	222209_s_at
cell cycle progression 1	CCPG1	9236	NP_065790	NM_020739	4D	222156_x_at
G proteincoupled receptor 89	GPR89	51463	NP_057418	NM_016334	4D	222140_s_at
Daspartate oxidase	DDO	8528	NP_004023	NM_004032	4D	222134_at
FtsJ homolog 2 (E. coli)	FTSJ2	29960	NP_037525	NM_013393	4D	222130_s_at
amphoterin induced gene 2	AMIGO2	347902	NP_862830	NM_181847	4D	222108_at
activating transcription factor 1	ATF1	466	NP_005162	NM_005171	4D	222103_at
solute carrier family 2 (facilitated glucose trans	SLC2A3	6515	NP_008862	NM_006931	4D	222088_s_at
interleukin 27 receptor, alpha	IL27RA	9466	NP_004834	NM_004843	4D	222062_at
hypothetical protein MGC3731	MGC3731	79159	NP_077289	NM_024313	4D	222057_at
zinc finger protein 45	ZNF45	7596	NP_003416	NM_003425	4D	222028_at

mannosidase, endoalpha						222817 at
	MANEA	79694	NP_078917	NM_024641	4D	222805_at
phosphotriesterase related	PTER	9317	NP_001001484	NM_001001484	4D	222798_at
PAP associated domain containing 1	PAPD1	55149	NP_060579	NM_018109	4D	222794_x_at
HSPC128 protein	HSPC128	29080	NP_054886	NM_014167	4D	222792_s_at
signal peptidase complex subunit 3 homolog (S cer	SPCS3	60559	NP_068747	NM_021928	4D	222753_s_at
sex comb on midleglike 1 (Drosophila)	SCML1	6322	NP_006737	NM_006746	4D	222747_s_at
chromosome 15 open reading frame 29	C15orf29	79768	NP_078989	NM_024713	4D	222745_s_at
trimethyllysine hydroxylase, epsilon	TMLHE	55217	NP_060666	NM_018196	4D	222744_s_at
hypothetical protein FLJ23375	FLJ23375	80021	NP_079232	NM_024956	4D	222739_at
hypothetical protein MGC5306	MGC5306	79101	NP_077021	NM_024116	4D	222728 s_at
lactamase, beta 2	LACTB2	51110	NP_057111	NM_016027	4D	222714_s_at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	222702_x_at
x 010 protein	MDS010	56983	NP_689518	NM_152305	4D	222681_at
RP42 homolog	RP42	54165	NP_065691	NM_020640	4D	222678_s_at
BAI1associated protein 2like 1	BAIAP2L1	55971	NP_061330	NM_018842	4D	222676_at
vmaf musculoaponeurotic fibrosarcoma oncogene homo	MAFB	9935	NP_005452	NM_005461	4D	222670_s_at
hypothetical protein FLJ11011	FLJ11011	55284	NP_001001481	NM_001001481	4D	222657_s_at
hypothetical protein FLJ11011	FLJ11011	55284	NP_001001481	NM_001001481	4D	222656_at
trichorhinophalangeal syndrome I	TRPS1	7227	NP_054831	NM_014112	4D	222651_s_at
exportin 4	XPO4	64328	NP_071904	NM_022459	4D	222649_at
hypothetical protein FLJ22662	FLJ22662	79887	NP_079105	NM_024829	4D	222639 s at
mediator of RNA polymerase II transcription, subun	MED28	80306	NP_079481	NM_025205	4D	222636_at
phosphatidylinositol 4kinase typeII beta	PI4K2B	55300	NP_060793	NM_018323	4D	222631_at
nudE nuclear distribution gene E homolog 1 (A. nid	NDEI	54820	NP_060138	NM_017668	4D	222625_s_at
chromosome 10 open reading frame 84	C10orf84	63877	NP_071346	NM_022063	4D	222617_s_at
exosome component l	EXOSCI	51013	NP_057130	NM_016046	4D	222609 s_at
xenotropic and polytropic retrovirus receptor	XPR1	9213	NP_004727	NM_004736	4D	222581_at
mitogenactivated protein kinase kinase kinase kina	MAP4K4	9448	NP_004825	NM_004834	4D	222547_at
zinc finger protein 395	ZNF395	55893	NP_061130	NM_018660	4D	222536_s_at
chromosome 14 open reading frame 108	C14orf108	55745	NP_060699	NM_018229	4D	222531_s_at

NM_014395 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_016060 NP_057144 51003 MED31 NM_02062 NP_075044 56919 DHX33 NM_022893 NP_075044 53335 BCL111A NM_022898 NP_075049 64919 BCL11B NM_022052 NP_075345 56000 NXF3 NM_022062 NP_071335 56000 NXF3 NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLNI NM_013438 NP_038466 29979 UBQLNI NM_016897 NP_057673 51300 C3orfi NM_0168997 NP_057673 51300 C3orfi NM_017830 NP_056997 51060 TLP19 NM_01685 NP_056997 51060 TLP19	HSPC182 protein	HSPC182	29101	NP_054907	NM_014188	4D	223053_x_at
NM_014395 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_0206162 NP_064547 56919 DHX33 NM_022893 NP_075049 56919 DHX33 NM_0214538 NP_075049 64919 BCL11A NM_0214538 NP_075049 64919 BCL11A NM_0205052 NP_075049 64919 BCL11B NM_021438 NP_055173 26253 CLEC4E NM_020981 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_057564 51247 PAIP2 NM_016480 NP_038466 29979 UBQLNI NM_015438 NP_038466 29979 UBQLNI NM_016897 NP_061870 54460 MRPS21 NM_017830 NP_057673 51300 C3orf1	solute carrier family 40 (ironregulated transporte	SLC40A1	30061	NP_055400	NM_014585	4D	223044_at
NM_012395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014396 NP_057044 51003 MED31 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014338 NP_075349 64919 BCL11B NM_014358 NP_071335 56000 NXF3 NM_014380 NP_08892 404636 FAM45A NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_018997 NP_061870 54460 MRPS21 NM_017830 NP_057673 51300 C3orf1 NM_015913 NP_066390 54940 OCIADI	Nacetyltransferase 5 (ARD1 homolog, S. cerevisiae)	NAT5	51126	NP_852669	NM_181528	4D	223040_at
NM_014395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_014360 NP_057144 51003 MED31 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_0753049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_01438 NP_066191 8708 B3CALT1 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_0156589 NP_057673 51300 C3orf1 NM_017830 NP_061870 54460 MRPS21 NM_015913 NP_066300 54940 OC1AD1	vacuolar protein sorting 29 (yeast)	VPS29	51699		NM_057180	4D	223026_s_at
NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_016480 NP_075734 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_015438 NP_038466 29979 UBQLN1 NM_015897 NP_061870 54460 MRPS21 NM_015899 NP_057673 51300 C3orf1 NM_015913 NP_060300 54460 MRPS21 NM_015913 NP_056997 51060 TLP19	likely ortholog of mouse nin one binding protein	NOBIP	28987	NP_054781	NM_014062	4D	223018_at
NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_020981 NP_071335 56000 NXF3 NM_020981 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_015438 NP_038466 29979 UBQLN1 NM_016589 NP_061870 54460 MRPS21 NM_016589 NP_057673 51300 C3orf1 NM_017830 NP_060300 54940 OCIAD1	endoplasmic reticulum thioredoxin superfamily memb	TLP19	51060	NP_056997	NM_015913	4D	223017_at
NM_012073 NP_071300 112399 EOLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022893 NP_075049 64919 BCL11B NM_014358 NP_075549 64919 BCL11B NM_014358 NP_075349 56000 NXF3 NM_020981 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_016589 NP_061870 54460 MRPS21 NM_032012 NP_114401 23731 C9orf5	OCIA domain containing 1	OCIADI	54940	NP_060300	NM_017830	4D	223011_s_at
NM_012395 NP_055210 27071 DAPPI NM_014395 NP_055210 27071 DAPPI NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_075049 64919 BCL11B NM_014358 NP_075049 64919 BCL11B NM_014358 NP_075049 64919 BCL11B NM_022052 NP_071335 56000 NXF3 NM_016480 NP_996892 404636 FAM45A NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_015897 NP_038466 29979 UBQLN1 NM_016899 NP_061870 54460 MRPS21 NM_01589 NP_057673 51300 C3orf1 NM_032012 NP_114401 23731 C9orf5	chromosome 9 open reading frame 5	C9orf5	23731	l'	NM_032012	4D	223008_s_at
NM_012303 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_0202052 NP_071335 56000 NXF3 NM_016480 NP_096892 404636 FAM45A NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLNI NM_013438 NP_038466 29979 UBQLNI NM_016899 NP_061870 54460 MRPS21 NM_016899 NP_057673 51300 C3orf1	chromosome 9 open reading frame 5	C9orf5	23731	l i	NM_032012	4D	223006_s_at
NM_022073 NF_07330 112393 EXLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_055210 27071 DAPP1 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_0202052 NP_071335 56000 NXF3 NM_016480 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_015997 NP_061870 54460 MRPS21	chromosome 3 open reading frame 1	C3orf1	51300		NM_016589	4D	223004 s_at
NM_0122073 NP_01330 112393 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_055210 27071 DAPP1 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_0751335 56000 NXF3 NM_022052 NP_071335 56000 NXF3 NM_016480 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_012479 NP_036611 7532 YWHAG NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1 NM_01505 NP_004996 4715 NDUFB9	mitochondrial ribosomal protein S21	MRPS21	54460	NP_061870	NM_018997	4D	222997_s_at
NM_022073 NP_07330 11239 EGLN3 NM_014395 NP_055210 27071 DAPPI NM_016060 NP_055210 27071 DAPPI NM_020162 NP_057144 51003 MED31 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11A NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_020981 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_013438 NP_038466 29979 UBQLN1 NM_013438 NP_038466 29979 UBQLN1	NADH dehydrogenase (ubiquinone) i beta subcomplex,	NDUFB9	4/15	NP_004996	NM_005005	4υ	222992_s_at
NM_0123073 NP_071330 112399 EGLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_016480 NP_08692 404636 FAM45A NM_016480 NP_057564 51247 PAIP2 NM_012479 NP_036611 7532 YWHAG NM_013438 NP_038466 29979 UBQLN1	ubiquilin 1	UBQLNI	29979	NP_038466	NM_013438	4D	222990_at
NM_022073 NP_071330 11239 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_020981 NP_066191 8708 B3GALT1 NM_016480 NP_057564 51247 PAIP2 NM_012479 NP_036611 7532 YWHAG	ubiquilin 1	UBQLNI	29979	NP_038466	NM_013438	4D	222989_s_at
NM_022073 NP_07330 112397 EGENS NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_020981 NP_996892 404636 FAM45A NM_016480 NP_057564 51247 PAIP2 NM_016480 NP_057564 51247 PAIP2	tyrosine 3monooxygenase/tryptophan 5monooxygenase	YWHAG	7532	NP_036611	NM_012479	4D	222985_at
NM_022073 NP_071330 112399 EGEN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_207009 NP_996892 404636 FAM45A NM_016480 NP_057564 51247 PAIP2	poly(A) binding protein interacting protein 2	PAIP2	51247	NP_057564	NM_016480	4D	222984_at
NM_022073 NP_071330 112399 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_02162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3 NM_020981 NP_066191 8708 B3GALT1	poly(A) binding protein interacting protein 2	PAIP2	51247	NP_057564	NM_016480	4D	222983_s_at
NM_022073 NP_071330 112399 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_079089 79871 FLJ13150 NM_014358 NP_075049 64919 BCL11B NM_022052 NP_071335 26253 CLEC4E NM_207009 NP_996892 404636 FAM45A	UDPGal:betaGlcNAc beta 1,3galactosyltransferase, p	B3GALT1	8708	NP_066191	NM_020981	4D	222969_at
NM_022073 NP_071330 112399 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_079089 79871 FLJ13150 NM_014358 NP_055173 26253 CLEC4E NM_022052 NP_071335 56000 NXF3	family with sequence similarity 45, member A	FAM45A	404636	NP_996892	NM_207009	4D	222955_s_at
INM_022073 INP_071330 I12393 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_075049 79871 FLJ13150 NM_014358 NP_055173 26253 CLEC4E	nuclear RNA export factor 3	NXF3	56000	NP_071335	NM_022052	4D	222949_at
NM_022073 NP_071330 112393 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_022898 NP_075049 64919 BCL11B	Ctype lectin domain family 4, member E	CLEC4E	26253	NP_055173	NM_014358	4D	222934_s_at
NM_022073 NP_071330 112393 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A NM_024813 NP_079089 79871 FLJ13150	Bcell CLL/lymphoma 11B (zinc finger protein)	BCL11B	64919	NP_075049	NM_022898	4D	222895_s_at
NM_022073 NP_071330 112393 EQLN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33 NM_022893 NP_075044 53335 BCL11A	hypothetical protein FLJ13150	FLJ13150	79871	NP_079089	NM_024813	4D	222893_s_at
NM_022073 NP_071330 112399 EGLN3 NM_014395 NP_055210 27071 DAPPI NM_016060 NP_057144 51003 MED31 NM_020162 NP_064547 56919 DHX33	Bcell CLL/lymphoma 11A (zinc finger protein)	BCLIIA	53335	NP_075044	NM_022893	4D	222891_s_at
NM_022073 NP_071330 112399 EGEN3 NM_014395 NP_055210 27071 DAPP1 NM_016060 NP_057144 51003 MED31	DEAH (AspGluAlaHis) box polypeptide 33	DHX33	56919	NP_064547	NM_020162	4D	222875_at
NM_014395 NP_055210 27071 DAPP1	mediator of RNA polymerase II transcription, subun	MED31	51003	NP_057144	NM_016060	4D	222867_s_at
NM_0220/3 NF_0/1330 112399 EQLINS	dual adaptor of phosphotyrosine and 3phosphoinosit	DAPPI	27071	NP_055210	NM_014395	4D	222858_s_at
מוא זמן מואר בפסבה זיויל	egl nine homolog 3 (C. elegans)	EGLN3	112399	NP_071356	NM_022073	4D	222847_s_at
D NM_198182 NP_937825 29841 TFCP2L2 transcription factor CP2like 2	transcription factor CP2like 2	TFCP2L2	29841	NP_937825	NM_198182	4D	222830_at

KIAA1333	KIAA1333	55632	NP_060239	NM_017769	4D	223256_at
KIAA1333	KIAA1333	55632	NP_060239	NM_017769	4D	223255_at
hypothetical protein MGC5309	MGC5309	84246	NP_115662	NM_032286	4D	223247_at
13kDa differentiationassociated protein	DAP13	55967	NP_061326	NM_018838	4D	223244_s_at
Fbox protein 8	FBXO8	26269	NP_036312	NM_012180	4D	223240_at
hypothetical protein DKFZp434K1421	DKFZP434K142	84081	NP_115517	NM_032141	4D	223236_at
cingulin	CGN	57530	NP_065821	NM_020770	4D	223233_s_at
ARV1 homolog (yeast)	ARVI	64801	NP_073623	NM_022786	4D	223223_at
zinc fingers and homeoboxes 1	ZHXI	11244	NP_009153	NM_007222	4D	223214_s_at
zinc finger, DHHC domain containing 16	ZDHHC16	84287	NP_932163	NM_198046	4D	223212_at
re						
SWI/SNFrelated, matrixassociated actindependent	SMARCADI	56916	NP_064544	NM_020159	4D	223197_s_at
sestrin 2	SESN2	83667	NP_113647	NM_031459	4D	223195_s_at
chromosome 6 open reading frame 69	C6orf69	222658	NP_775833	NM_173562	4D	223176_at
ubiquitin specific protease 25	USP25	29761	NP_037528	NM_013396	4D	223167_s_at
zinc finger, C3HC type 1	ZC3HC1	51530	NP_057562	NM_016478	4D	223163_s_at
chromosome 4 open reading frame 14	C4orf14	84273	NP_115689	NM_032313	4D	223157_at
hypothetical protein MGC2714	MGC2714	84259	NP_115675	NM_032299	4D	223151_at
DEAH (AspGluAlaHis) box polypeptide 36	DHX36	170506	NP_065916	NM_020865	4D	223139_s_at
androgeninduced 1	AIGI	51390	NP_057192	NM_016108	4D	223136_at
hypothetical protein H17	H17	55572	NP_060017	NM_017547	4D	223128_at
secreted frizzledrelated protein 2	SFRP2	6423	NP_003004	NM_003013	4D	223121_s_at
hypothetical protein HSPC196	HSPC196	51524	NP_057548	NM_016464	4D	223113_at
ADPribosylhydrolase like 2	ADPRHL2	54936	NP_060295	NM_017825	4D	223097_at
mitochondrial ribosomal protein L51	MRPL51	51258	NP_057581	NM_016497	4D	223086_x_at
ring finger protein 19	RNF19	25897	NP_056250	NM_015435	4D	223085_at
cyclin Dtype bindingprotein 1	CCNDBP1	23582	NP_036274	NM_012142	4D	223084_s_at
hypothetical protein MGC2941	MGC2941	79142	NP_077273	NM_024297	4D	223081_at
NOL1/NOP2/Sun domain family, member 2	NSUN2	54888	NP_060225	NM_017755	4D	223076_s_at
hypothetical protein FLJ14753	FLJ14753	84641	NP_115947	NM_032558	4D	223073_at
selenoprotein K	SELK	58515	NP_067060	NM_021237	4D	223070_at
STARD3 Nterminal like	STARD3NL	83930	NP_114405	NM_032016	4D	223065_s_at
hypothetical protein LOC51255	LOC51255	51255	NP_057578	NM_016494	4D	223064_at
exportin 5	XPO5	57510	NP_065801	NM_020750	4D	223057_s_at

	HAGHL	84264	NP 115680	NM_032304	4D	223485 at
homo	HIMIMIOD	14004	INI_020271	INIM_O12407	ţ	10_01+C24
tranclogge of inner mitochondrial membrane	TIMM8R	10590	ND 036591	NM 012459		223478 at
phosphatidylinositol glycan, class M	PIGM	93183	NP_660150	NM_145167	4D	223470 at
calcium/calmodulindependent protein kinase kinase	CAMKKI	84254	NP_115670	NM_032294	4D	223460_at
peroxisome proliferative activated receptor, alpha	PPARA	5465	NP_001001928	NM_001001928	4D	223437_at
chromosome 7 open reading frame 36	C7orf36	57002	NP_064577	NM_020192	4D	223433_at
cappuccino	CNO	55330	NP_060836	NM_018366	4D	223431_at
hypothetical protein FLJ20425	LYAR	55646	NP_060286	NM_017816	4D	223413_s_at
kelch repeat and BTB (POZ) domain containing	KBTBD7	84078	NP_115514	NM_032138	4D	223412_at
chromosome 7 open reading frame 35	C7orf35	85025	NP_116325	NM_032936	4D	223396_at
SERTA domain containing 1	SERTADI	29950	NP_037508	NM_013376	4D	223394_at
zinc finger protein 537	ZNF537	57616	NP_065907	NM_020856	4D	223393_s_at
zinc finger protein 537	ZNF537	57616	NP_065907	NM_020856	4D	223392_s_at
tripartite motifcontaining 4	TRIM4	89122	NP_149082	NM_033091	4D	223384_s_at
pleckstrin homology domain containing, family A (p	PLEKHA3	65977	NP_061964	NM_019091	4D	223370_at
melanoma associated antigen (mutated) l	MUMI	84939	NP_116242	NM_032853	4D	223347_at
histone deacetylase 8	HDAC8	55869	NP_060956	NM_018486	4D	223345_at
ribonucleotide reductase M2 B (TP53 inducible)	RRM2B	50484	NP_056528	NM_015713	4D	223342_at
ATPase inhibitory factor 1	ATPIF1	93974	NP_835497	NM_178190	4D	223338_s_at
serologically defined colon cancer antigen 10	SDCCAG10	10283	NP_005860	NM_005869	4D	223337_at
RAB18, member RAS oncogene family	RAB18	22931	NP_067075	NM_021252	4D	223336_s_at
2 hypothetical protein DKFZp586C1924	DKFZp586C192	84233	NP_115649	NM_032273	4D	223334_at
hypothetical protein FLJ23518	FLJ23518	79780	NP_079001	NM_024725	4D	223301_s_at
5'nucleotidase, cytosolic III	NT5C3	51251	NP_057573	NM_016489	4D	223298_s_at
hypothetical protein MGC4268	MGC4268	83607	NP_113633	NM_031445	4D	223297_at
mitochondrial carrier protein	MGC4399	84275	NP_115691	NM_032315	4D	223296_at
chromosome X open reading frame 26	CXorf26	51260	NP_057584	NM_016500	4D	223294_at
K562 cellderived leucinezipperlike protein 1	KLPI	57106	NP_065111	NM_020378	4D	223284_at
putative small membrane protein NID67	NID67	85027	NP_116565	NM_032947	4D	223276_at
PTD012 protein	PTD012	28970	NP_054758	NM_014039	4D	223268_at
ORM1like 3 (S. cerevisiae)	ORMDL3	94103	NP_644809	NM_139280	4D	223259_at

regulator of Gprotein signalling 18	RGS18	64407	NP_570138	NM_130782	4D	223809_at
GTP binding protein 2	GTPBP2	54676	NP_061969	NM_019096	4D	223789_s_at
kinesin family member 9	KIF9	64147	NP_071737	NM_022342	4D	223778_at
dystrobrevin binding protein 1	DTNBPI	84062	NP_898861	NM_183040	4D	223763_at
general transcription factor IIH, polypeptide 2, 4	GTF2H2	2966	NP_001506	NM_001515	4D	223758_s_at
hypothetical protein MGC13057	MGC13057	84281	NP_115697	NM_032321	4D	223754_at
C1q and tumor necrosis factor related protein 2	C1QTNF2	114898	NP_114114	NM_031908	4D	223749_at
mitochondrial ribosomal protein L4	MRPL4	51073	NP_057040	NM_015956	4D	223743_s_at
thymocyte protein thy28	THY28	29087	NP_054893	NM_014174	4D	223711_s_at
winglesstype MMTV integration site family, member	WNT10A	80326	NP_079492	NM_025216	4D	223709_s_at
chromosome 22 open reading frame 23	C22orf23	84645	NP_115950	NM_032561	4D	223706_at
latent transforming growth factor beta binding pro	LTBP2	4053	NP_000419	NM_000428	4D	223690_at
hypothetical protein MGC11102	MGC11102	84285	NP_115701	NM_032325	4D	223682_s_at
CDC42 small effector 1	CDC42SE1	56882	NP_064624	NM_020239	4D	223674_s_at
CGI30 protein	CG130	51611	NP_057042	NM_015958	4D	223671_x_at
hemogen	HEMGN	55363	NP_060907	NM_018437	4D	223670_s_at
nuclear receptor binding factor 2	NRBF2	29982	NP_110386	NM_030759	4D	223650_s_at
Jtype cochaperone HSC20	HSC20	150274	NP_741999	NM_172002	4D	223647_x_at
zinc ribbon domain containing, 1	ZNRD1	30834	NP_055411	NM_014596	4D	223639_s_at
RASD family, member 2	RASD2	23551	NP_055125	NM_014310	4D	223634_at
family with sequence similarity 14, member A	FAM14A	83982	NP_114425	NM_032036	4D	223626_x_at
G proteincoupled receptor 34	GPR34	2857	NP_005291	NM_005300	4D	223620_at
ropporin 1like	ROPNIL	83853	NP_114122	NM_031916	4D	223609_at
kelch repeat and BTB (POZ) domain containing 2	KBTBD2	25948	NP_056298	NM_015483	4D	223584_s_at
hypothetical protein FLJ23467	FLJ23467	79626	NP_078851	NM_024575	4D	223583_at
chromosome 6 open reading frame 203	C6orf203	51250	NP_057571	NM_016487	4D	223576_at
chromosome 1 open reading frame 26	Clorf26	54823	NP_060143	NM_017673	4D	223548_at
ankyrin repeat domain 32	ANKRD32	84250	NP_115666	NM_032290	4D	223542_at
nudix (nucleoside diphosphate linked moiety X)type	NUDI 12	83394	NP_113626	NM_031438	40	223535_at
G proteincoupled receptor 89	GPR89	51463	NP_057418	NM_016334	4D	22353 L_x_at
FLJ20859 gene	FLJ20859	64745	NP_073571	NM_022734	4D	223528_s_at
chromosome 9 open reading frame 45	C9orf45	81571	NP_110441	NM_030814	4D	223522_at
Rasassociated protein Rap1	RBJ	51277	NP_057628	NM_016544	4D	223504_at

171 A 1787	KIAA1287	57508	NP 065799	NM 020748	4D	224308 s at
SECIS binding protein 2	SECISBP2	79048	NP_076982	NM_024077	4D	224250_s_at
Fas (TNFRSF6) associated factor 1	FAF1	11124	NP_008982	NM_007051	4D	224217_s_at
CG130 protein	CG130	51611	NP_057042	NM_015958	4D	224196_x_at
heat shock 70kDa protein 8	HSPA8	3312	NP_006588	NM_006597	4D	224187_x_at
chromosome X open reading frame 26	CXorf26	51260	NP_057584	NM_016500	4D	224177_s_at
tripartite motifcontaining 4	TRIM4	89122	NP_148977	NM_033017	4D	224159_x_at
interleukin 17 receptor B	IL17RB	55540	NP_061195	NM_018725	4D	224156_x_at
hypothetical protein MGC10561	MGC10561	84733	NP_116036	NM_032647	4D	224138_at
calcium channel, voltagedependent, gamma subunit 7	CACNG7	59284	NP_114102	NM_031896	4D	224137_at
G proteincoupled receptor 81	GPR81	27198	NP_115943	NM_032554	4D	224131_at
potassium voltagegated channel, subfamily H (eagre	KCNH7	90134	NP_150375	NM_033272	4D	224099_at
hippocampus abundant gene transcriptlike 2	HIATL2	84278	NP_115694	NM_032318	4D	224078_at
visual system homeobox 1 homolog, CHX10like (zebra	VSXI	30813	NP_955457	NM_199425	4D	224074_at
zinc finger, CW type with PWWP domain 1	ZCWPW1	55063	NP_060454	NM_017984	4D	223992_x_at
FLJ20859 gene	FLJ20859	64745	NP_073571	NM_022734	4D	223988_x_at
IGFII mRNAbinding protein 2	IMP2	10644	NP_006539	NM_006548	4D	223963_s_at
transmembrane protease, serine 3	TMPRSS3	64699	NP_115781	NM_032405	4D	223949_at
transmembrane protease, serine 3	TMPRSS3	64699	NP_076927	NM_024022	4D	223948_s_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST5	23563		NM_012126	4D	223942_x_at
solute carrier family 39 (zinc transporter), membe	SLC39A3	29985	NP_653165	NM_144564	4D	223917_s_at
BCL6 corepressor	BCOR	54880	NP_065977	NM_020926	4D	223915_at
epsin 3	EPN3	55040	NP_060427	NM_017957	4D	223895_s_at
G proteincoupled receptor 132	GPR132	29933	NP_037477	NM_013345	4D	223887_at
chromosome 20 open reading frame 24	C20orf24	55969	NP_061328	NM_018840	4D	223880_x_at
armadillo repeat containing 2	ARMC2	84071	NP_115507	NM_032131	4D	223866_at
hypothetical protein LOC51234	LOC51234	51234	NP_057538	NM_016454	4D	223857_x_at
orthopedia homolog (Drosophila)	OTP	23440	NP_115485	NM_032109	4D	223835_x_at
programmed cell death 1 ligand 1	PDCD1LG1	29126	NP_054862	NM_014143	4D	223834_at
KIAA1160 protein	KIAA1160	57461	NP_065752	NM_020701	4D	223831_x_at
hypothetical protein radiovoz	1 10000					

CDW92 antigen	CDW92	23446	NP_536856	NM_080546	4D	224595_at
zinc finger protein ZFOC1	ZFOC1	144348	NP_689650	NM_152437	4D	224593_at
chromosome 20 open reading frame 30	C20orf30	29058	NP_054864	NM_014145	4D	224584_at
chromosome 3 open reading frame 10	C3orf10	55845	NP_060932	NM_018462	4D	224575_at
reticulon 3	RTN3	10313	NP_958831	NM_201428	4D	224564_s_at
WAS protein family, member 2	WASF2	10163	NP_008921	NM_006990	4D	224563_at
murine retrovirus integration site 1 homolog	MRVII	10335	NP_006060	NM_006069	4D	224550_s_at
CXXC finger 5	CXXC5	51523	NP_057547	NM_016463	4D	224516_s_at
thioredoxinlike 5	TXNL5	84817	NP_116120	NM_032731	4D	224511_s_at
hypothetical protein MGC10744	MGC10744	84314	NP_898888	NM_183065	4D	224496_s_at
hypothetical protein MGC10744	MGC10744	84314	NP_898888	NM_183065	4D	224495_at
HECT domain containing 1	HECTDI	25831	NP_056197	NM_015382	4D	224481_s_at
leucine zipper, putative tumor suppressor 2	LZTS2	84445	NP_115805	NM_032429	4D	224473_x_at
PYM protein	PYM	84305	NP_115721	NM_032345	4D	224465_s_at
chromosome 9 open reading frame 125	C9orf125	84302	NP_115718	NM_032342	4D	224458_at
hypothetical protein MGC12966	MGC12966	84792	NP_116095	NM_032706	4D	224452_s_at
RIO kinase 1 (yeast)	RIOK1	83732	NP_113668	NM_031480	4D	224450_s_at
ring finger protein 7	RNF7	9616	NP_899060	NM_183237	4D	224439_x_at
chromosome 10 open reading frame 58	C10orf58	84293	NP_115709	NM_032333	4D	224435_at
mitochondrial translation optimization 1 homolog (MTO1	25821	NP_036255	NM_012123	4D	224430_s_at
mediator of RNA polymerase II transcription, subun	MED28	80306	NP_079481	NM_025205	4D	224416_s_at
histidine triad nucleotide binding protein 2	HINT2	84681	NP_115982	NM_032593	4D	224415_s_at
ring finger protein 7	RNF7	9616	NP_055060	NM_014245	4D	224395_s_at
ring finger protein 17	RNF17	56163	NP_114383	NM_031994	4D	224383_at
TAF7like RNA polymerase II, TATA box binding prote	TAF7L	54457	NP_079161	NM_024885	4D	224380_s_at
RAB18, member RAS oncogene family	RAB18	22931	NP_067075	NM_021252	4D	224377_s_at
NDRG family member 3	NDRG3	57446	NP_114402	NM_032013	4D	224368_s_at
ubiquitinconjugating enzyme E2, J2 (UBC6 homolog,	UBE2J2	118424	NP_919297	NM_194316	4D	224347_x_at
mitochondrial ribosomal protein L51	MRPL51	51258	NP_057581	NM_016497	4D	224334_s_at
mitochondrial ribosomal protein L36	MRPL36	64979	NP_115868	NM_032479	4D	224331_s_at
mitochondrial ribosomal protein L27	MRPL27	51264	NP_683412	NM_148571	4D	224330_s_at

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chromosome 9 open reading frame 123	C9orf123	90871	XP_372075	XM_372075	4D	224879_at
chromosome 9 open reading frame 123	C9orf123	90871	XP_372075	XM_372075	4D	224860_at
tumor protein p53 inducible nuclear protein 2	TP53INP2	58476	NP_067025	NM_021202	4D	224836_at
cytoplasmic polyadenylation element binding protei	CPEB4	80315	NP_085130	NM_030627	4D	224831_at
cytoplasmic polyadenylation element binding protei	CPEB4	80315	NP_085130	NM_030627	4D	224828_at
family with sequence similarity 36, member A	FAM36A	116228	NP_932342	NM_198076	4D	224824_at
family with sequence similarity 36, member A	FAM36A	116228	NP_932342	NM_198076	4D	224820_at
transcription elongation factor A (SII)like 8	TCEAL8	90843	NP_699164	NM_153333	4D	224819_at
3hydroxyisobutyrate dehydrogenase	HIBADH	111112	NP_689953	NM_152740	4D	224812_at
arrestin domain containing 3	ARRDC3	57561	NP_065852	NM_020801	4D	224797_at
RAB18, member RAS oncogene family	RAB18	22931	NP_067075	NM_021252	4D	224787_s_at
hypothetical protein MGC29814	MGC29814	283991	NP_872371	NM_182565	4D	224785_at
RNA binding motif protein 17	RBM17	84991	NP_116294	NM_032905	4D	224780_at
hypothetical protein FLJ22875	FLJ22875	84191	NP_115607	NM_032231	4D	224779_s_at
hypothetical protein FLJ10006	FLJ10006	55677	NP_060439	NM_017969	4D	224768_at
pim3 oncogene	PIM3	415116	NP_001001852	NM_001001852	4D	224739_at
cell division cycle and apoptosis regulator 1	CCAR1	55749	NP_060707	NM_018237	4D	224736_at
hypothetical protein FLJ12519	FLJ12519	84128	NP_115544	NM_032168	4D	224721_at
likely ortholog of mouse gene rich cluster, C10 ge	GRCC10	113246	NP_612434	NM_138425	4D	224719_s_at
MKI67 (FHA domain) interacting nucleolar phosphopr	MKI67IP	84365	NP_115766	NM_032390	4D	224713_at
chromosome 20 open reading frame 35	C20orf35	55861	NP_291020	NM_033542	4D	224669_at
kinesin family member 5B	KIF5B	3799	NP_004512	NM_004521	4D	224662_at
hypothetical protein MGC14156	MGC14156	84992	NP_116295	NM_032906	4D	224661_at
mitogeninducible gene 6	MIG6	54206	NP_061821	NM_018948	4D	224657_at
adenylate kinase 3 like 1	AK3L1	50808	NP_057366	NM_016282	4D	224655_at
signal peptide peptidase 3	SPPL3	121665	NP_620584	NM_139015	4D	224640_at
similar to RIKEN cDNA 2310016E02	LOC400948	400948	XP_376043	XM_376043	4D	224637_at
G patch domain containing 4	GPATC4	54865	NP_056405	NM_015590	4D	224634_at
G patch domain containing 4	GPATC4	54865	NP_872620	NM_182679	4D	224632_at
small EDRKrich factor 2	SERF2	10169	NP_005761	NM_005770	4D	224625_x_at
dynein, cytoplasmic, light intermediate polypeptid	DNCLI2	1783	NP_006132	NM_006141	4D	224614_at
C1L2 gene	CTL2	57153	NP_065161	NM_020428	4D	224609_at

protein phosphatase 1, regulatory (inhibitor) subu Rho GTPase activating protein 18 RAB11 family interacting protein 1 (class I) tetratricopeptide repeat domain 14 chromosome 21 open reading frame 4	C21orf4	757	NP_006125	NM_000134		225162
8	***************************************	131613			4	225182_at
8	TTC14		NP_597719			11
	RABITFIP	80223	NP_001002814	NM_001002814		225177_at
protein phosphatase 1, regulatory (inhibitor) subu	ARHGAP18	93663	NP_277050	NM_033515	4D	225173_at
FERM domain containing 4A	PPPIRIB	84152	NP_852606	NM_181505	4D	225165_at
PERM 1	FRMD4A	55691	NP_060497	NM_018027	4D	225163_at
hypothetical protein MGC4549	MGC4549	84337	NP_115753	NM_032377	4D	225156_at
(Droso			esta-			2000
synapse associated protein 1. SAP47 homolog	SYAPI	94056	NP_116185	NM_032796	4D	225154_at
G elongation factor, mitochondrial 1	GFM1	85476	NP_079272	NM_024996	4D	225153_at
	KIAA1718	80853	XP_376680	XM_376680	4D	225142_at
Kruppellike factor 3 (basic)	KLF3	51274	NP_057615	NM_016531	4D	225133_at
rabaptin, RAB GTPase binding effector protein 1	RABEPI	9135	NP_004694	NM_004703	4D	225092_at
synovial apoptosis inhibitor 1, synoviolin	INAAS	84447	NP_115807	NM_032431	4D	225090_at
hypothetical protein FLJ31153	FLJ31153	123811	NP_653201	NM_144600	4D	225088_at
cleavage and polyadenylation specific factor 3, 73	CPSF3	51692	NP_057291	NM_016207	4D	225082_at
RAB2B, member RAS oncogene family	RAB2B	84932	NP_116235	NM_032846	4D	225074_at
chromosome 6 open reading frame 68	C6orf68	116150	NP_612468	NM_138459	4D	225071_at
DnaJ (Hsp40) homolog, subfamily A, member 4	DNAJA4	55466	NP_061072	NM_018602	4D	225061_at
angiotensin II receptorassociated protein	AGTRAP	57085	NP_065083	NM_020350	4D	225059_at
zinc finger protein 512	ZNF512	84450	NP_115810	NM_032434	4D	225050_at
solute carrier family 15, member 4	SLC15A4	121260	NP_663623	NM_145648	4D	225043_at
chromosome 9 open reading frame 105	C9orf105	401505	NP_001001790	NM_001001790	4D	225036_at
chromodomain helicase DNA binding protein 6	CHD6	84181	NP_115597	NM_032221	4D	225031_at
hypothetical gene supported by BC032431	LOC389203	389203	XP_371691	XM_371691	4D	225014_at
ataxin 7like 3	ATXN7L3	56970	XP_375456	XM_375456	4D	224969_at
solute carrier family 26 (sulfate transporter), me	SLC26A2	1836	NP_000103	NM_000112	4D	224963_at
solute carrier family 26 (sulfate transporter), me	SLC26A2	1836	NP_000103	NM_000112	4D	224959_at
82kD FMRP Interacting Protein	182FIP	57532	NP_065823	NM_020772	4D	224956_at
mitochondrial ribosomal protein S24	MRPS24	64951	NP_114403	NM_032014	4D	224948_at
chromosome 22 open reading frame 16	C22orf16	400916	NP_998885	NM_213720	4D	224932_at
myeloidassociated differentiation marker	MYADM	91663	NP_612382	NM_138373	4D	224920_x_at
translocase of inner mitochondrial membrane 50 hom	TIMM50	92609	NP_001001563	NM_001001563	4D	224913_s_at

translocase of inner mitochondrial membrane 23 hom	TIMM23	10431	NP_006318	NM_006327	4D	225535_s_at
CCAAT/enhancer binding protein (C/EBP), gamma	CEBPG	1054	NP_001797	NM_001806	4D	225527_at
chromosome 20 open reading frame 81	C20orf81	64773	NP_073597	NM_022760	4D	225505_s_at
transmembrane protein 18	TMEM18	129787	NP_690047	NM_152834	4D	225489_at
mesoderm induction early response 1	MIER1	57708	NP_065999	NM_020948	4D	225475_at
nucleoporin 35kDa	NUP35	129401	NP_612142	NM_138285	4D	225470_at
glycerol3phosphate acyltransferase, mitochondrial	GPAM	57678	NP_065969	NM_020918	4D	225424_at
glycerol3phosphate acyltransferase, mitochondrial	GPAM	57678	NP_065969	NM_020918	4D	225420_at
ring finger protein 149	RNF149	284996	NP_775918	NM_173647	4D	225414_at
chromosome I open reading frame 19	Clorf19	116461	NP_443197	NM_052965	4D	225399_at
dedicator of cytokinesis 7	DOCK7	85440	NP_212132	NM_033407	4D	225384_at
endothelial cell adhesion molecule	ESAM	90952	NP_620411	NM_138961	4D	225369_at
hypothetical protein PP2447	PP2447	80305	NP_079480	NM_025204	4D	225360_at
homolog of yeast TIM14	TIM14	131118	NP_957713	NM_201261	4D	225359_at
chromosome 3 open reading frame 6	C3orf6	152137	NP_848018	NM_178335	4D	225331_at
acylCoenzyme A binding domain containing 6	ACBD6	84320	NP_115736	NM_032360	4D	225317_at
COMM domain containing 6	COMMD6	170622	NP_987091	NM_203495	4D	225312_at
polyribonucleotide nucleotidyltransferase 1	PNPTI	87178	NP_149100	NM_033109	4D	225291_at
hypothetical protein AL133206	LOC64744	64744	NP_073570	NM_022733	4D	225282_at
spermidine/spermine N l acetyltransferase 2	SAT2	112483	NP_597998	NM_133491	4D	225272_at
TH1like (Drosophila)	THIL	51497	NP_945327	NM_198976	4D	225261_x_at
mitochondrial ribosomal protein L32	MRPL32	64983	NP_114109	NM_031903	4D	225260_s_at
mitochondrial ribosomal protein L35	MRPL35	51318	NP_663619	NM_145644	4D	225255_at
sarcolemma associated protein	SLMAP	7871	NP_009090	NM_007159	4D	225243_s_at
musashi homolog 2 (Drosophila)	MSI2	124540	NP_620412	NM_138962	4D	225237_s_at
hypothetical LOC83640	MGC2560	83640	NP_113640	NM_031452	4D	225210_s_at
mitochondrial ribosomal protein L14	MRPL14	64928	NP_115487	NM_032111	4D	225201_s_at
mitochondrial ribosomal protein S26	MRPS26	64949	NP_110438	NM_030811	4D	225196_s_at
zinc finger, CSL domain containing 2	ZCSL2	285381	NP_996662	NM_206831	4D	225195_at
Ras association (RalGDS/AF6) and pleckstrin homolo	RAPHI	65059	NP_998754	NM_213589	4D	225189_s_at
Ras association (RalGDS/AF6) and pleckstrin homolo	RAPH1	65059	NP_998754	NM_213589	4D	225188_at

solute carrier family 35, member B4	SLC35B4	84912	NP_116215	NM_032826	4D	225881_at
hypothetical protein dJ462O23.2	DJ462O23.2	57185	NP_065181	NM_020448	4D	225876_at
solute carrier family 35, member F5	SLC35F5	80255	NP_079457	NM_025181	4D	225872_at
glucosaminephosphate Nacetyltransferase 1	GNPNATI	64841	NP_932332	NM_198066	4D	225853_at
ankyrin repeat domain 17	ANKRD17	26057	NP_942592	NM_198889	4D	225852_at
chromosome 6 open reading frame 83	C6orf83	113402	NP_660152	NM_145169	4D	225850_at
chromosome 6 open reading frame 83	C6orf83	113402	NP_660152	NM_145169	4D	225849_s_at
hypothetical protein MGC13204	MGC13204	83695	NP_113653	NM_031465	4D	225837_at
hypothetical protein MGC13204	MGC13204	83695	NP_113653	NM_031465	4D	225836_s_at
hypothetical gene supported by AL449243	LOC91689	91689	NP_201575	NM_033318	4D	225795_at
methionine sulfoxide reductase B3	MSRB3	253827	NP_932346	NM_198080	4D	225790_at
methionine sulfoxide reductase B3	MSRB3	253827	NP_932346	NM_198080	4D	225782_at
chromosome 6 open reading frame 93	C6orf93	84946	NP_116249	NM_032860	4D	225748_at
zinc finger, DHHC domain containing 8	ZDHHC8	29801	NP_037505	NM_013373	4D	225744_at
RNA pseudouridylate synthase domain containing	RPUSD3	285367	NP_775930	NM_173659	4D	225743_at
THUMP domain containing 3	THUMPD3	25917	NP_056268	NM_015453	4D	225741_at
mitochondrial ribosomal protein L55	MRPL55	128308	NP_852129	NM_181464	4D	225719_s_at
KIAA1715	KIAA1715	80856	NP_085153	NM_030650	4D	225718_at
hypothetical protein BC009518	LOC90799	90799	NP_612372	NM_138363	4D	225705_at
TIGAI	TIGAI	114915	NP_443726	NM_053000	4D	225698_at
family with sequence similarity 33, member A	FAM33A	348235	NP_872426	NM_182620	4D	225686_at
interferon (alpha, beta and omega) receptor 1	IFNARI	3454	NP_000620	NM_000629	4D	225661_at
hypothetical protein LOC339745	LOC339745	339745	NP_001001664	NM_001001664	4D	225659_at
ubiquitinconjugating enzyme E2E 2 (UBC4/5 homolog,	UBE2E2	7325	NP_689866	NM_152653	4D	225651_at
similar to hypothetical protein 9530023G02	MGC90512	121642	NP_001001655	NM_001001655	4D	225625_at
beta1,3Nacetylglucosaminyltransf						
UDPGlcNAc:betaGal	B3GNT5	84002	NP_114436	NM_032047	4D	225612_s_at
hypothetical protein FLJ31795	FLJ31795	124808	NP_653210	NM_144609	4D	225607_at
chromosome 9 open reading frame 19	C9orf19	152007	NP_071738	NM_022343	4D	225604_s_at
SH3 multiple domains 2	SH3MD2	57630	NP_065921	NM_020870	4D	225589_at
putative acylCoA dehydrogenase	FLJ12592	84129	NP_115545	NM_032169	4D	225573_at
chromosome 3 open reading frame 19	C3orf19	51244	NP_057558	NM_016474	4D	225559_at
zinc finger, CCHC domain containing 9	ZCCHC9	84240	NP_115656	NM_032280	4D	225538_at

hynothetical protein LOC 116068	LOC116068	116068	XP_371760	XM 371760	4D	226321_at
THO complex 4	THOC4	10189	NP_005773	NM_005782	4D	226319_s_at
chromosome 6 open reading frame 192	C6orf192	116843	NP_439896	NM_052831	4D	226301_at
mitochondrial ribosomal protein S15	MRPS15	64960	NP_112570	NM_031280	4D	226296_s_at
hypothetical protein MGC23909	MGC23909	153339	NP_777569	NM_174909	4D	226276_at
jun dimerization protein 2	JDP2	122953	NP_569736	NM_130469	4D	226267_at
fastinginducible integral membrane protein TM6P1	FLJ90024	129303	NP_699173	NM_153342	4D	226239_at
chromosome 13 open reading frame 8	C13orf8	283489	NP_115812	NM_032436	4D	226194_at
glycolipid transfer protein	GLTP	51228	NP_057517	NM_016433	4D	226177_at
hypothetical protein MGC29649	MGC29649	283237	NP_776171	NM_173810	4D	226175_at
SET binding factor 2	SBF2	81846	NP_112224	NM_030962	4D	226169_at
hypothetical protein LOC285636	LOC285636	285636	NP_787117	NM_175921	4D	226159_at
retinoic acid induced 1	RAH	10743	NP_109590	NM_030665	4D	226143_at
hypothetical protein MGC23280	MGC23280	147015	NP_653284	NM_144683	4D	226121_at
TRAFinteracting protein with a forkheadassociated	TIFA	92610	NP_443096	NM_052864	4D	226117_at
fibronectin type III domain containing 5	FNDC5	252995	NP_715637	NM_153756	4D	226097_at
membrane protein, palmitoylated 5 (MAGUK p55 subfa	MPP5	64398	NP_071919	NM_022474	4D	226092_at
synaptotagmin XIII	SYT13	57586	NP_065877	NM_020826	4D	226086_at
caspase 2, apoptosisrelated cysteine protease (neu	CASP2	835	NP_116764	NM_032982	4D	226036_x_at
disrupted in renal carcinoma 2	DIRC2	84925	NP_116228	NM_032839	4D	226026_at
copper metabolism (Murr1) domain containing 1	COMMD1	150684	NP_689729	NM_152516	4D	226024_at
hypothetical protein Ells1	Ells1	222166	NP_690006	NM_152793	4D	226018_at
coiledcoil domain containing 12	CCDC12	151903	NP_653317	NM_144716	4D	226011_at
HESB like domain containing 1	НВГОІ	122961	NP_919255	NM_194279	4D	226007_at
cleavage and polyadenylation specific factor 2, 10	CPSF2	53981	NP_059133	NM_017437	4D	225994_at
hypothetical protein MGC15677	MGC15677	84964	NP_116267	NM_032878	4D	225969_at
adult retina protein	LOC153222	153222	NP_705835	NM_153607	4D	225957_at
zinc finger protein 655	ZNF655	79027	NP_612503	NM_138494	4D	225945_at
CREBBP/EP300 inhibitor 2	CRI2	163126	NP_694964	NM_153232	4D	225936_at
FLJ45445 protein	FLJ45445	399844	NP_001004321	NM_001004321	4D	225899_x_at
hypothetical protein FLJ13089	FLJ13089	81008	NP_079229	NM_024953	4D	225888_at
solute carrier family 35, member B4	SLC35B4	84912	NP_116215	NM_032826	4D	225882_at

hypothetical protein LOC152485	LOC152485	152485	NP_849157	NM_178835	40	226764_at
mitochondrial ribosomal protein S9	MRPS9	64965	NP_872578	NM_182640	4D	226749_at
hypothetical protein KIAA1991	KIAA1991	254225	XP_495886	XM_495886	4D	226739_at
2 hypothetical protein DKFZp313N0621	DKFZp313N062	285343	NP_776187	NM_173826	4D	226688_at
solute carrier family 26, member 11	SLC26A11	284129	NP_775897	NM_173626	4D	226679_at
cartilage associated protein	CRTAP	10491	NP_006362	NM_006371	4D	226656_at
NudC domain containing 2	NUDCD2	134492	NP_660309	NM_145266	4D	226642_s_at
hypothetical protein MGC5391	MGC5391	84826	NP_116129	NM_032740	4D	226639_at
phospholipase D1, phophatidylcholinespecific	PLD1	5337	NP_002653	NM_002662	4D	226636_at
SUMO1/sentrin specific protease 1	SENP1	29843	NP_055369	NM_014554	4D	226619_at
NADH dehydrogenase (ubiquinone) flavoprotein 3, 10	NDUFV3	4731	NP_066553	NM_021075	4D	226616_s_at
chromosome 8 open reading frame 13	C8orf13	83648	NP_444509	NM_053279	4D	226614_s_at
proline rich 6	PRR6	201161	NP_859067	NM_181716	4D	226611_s_at
chromosome 7 open reading frame 6	C7orf6	219285	NP_689916	NM_152703	4D	226603_at
zinc finger protein 406	ZNF406	57623	NP_065914	NM_020863	4D	226564_at
histidine triad nucleotide binding protein 3	HINT3	135114	NP_612638	NM_138571	4D	226537_at
hypothetical protein FLJ14466	FLJ14466	84876	NP_116179	NM_032790	4D	226531_at
hypothetical protein MGC26717	MGC26717	285237	NP_776185	NM_173824	4D	226524_at
similar to RIKEN cDNA 0610011N22	LOC133957	133957	NP_660308	NM_145265	4D	226515_at
p21/Cdc42/Rac1activated kinase 1 (STE20 homolog, y	PAKI	5058	NP_002567	NM_002576	4D	226507_at
transmembrane and coiledcoil domains 3	TMCC3	57458	NP_065749	NM_020698	4D	226489_at
hypothetical protein FLJ14721	FLJ14721	84915	NP_116218	NM_032829	4D	226487_at
gammaglutamyltransferaselike 3	GGTL3	2686	NP_439895	NM_052830	4D	226470_at
chromosome 9 open reading frame 42	C9orf42	116224	NP_612206	NM_138333	4D	226443_at
ankyrin repeat and BTB (POZ) domain containing	ABTBI	80325	NP_115937	NM_032548	4D	226442_at
PTX1 protein	PTXI	51290	NP_057654	NM_016570	4D	226422_at
membraneassociated ring finger (C3HC4) 5	MARCH5	54708	NP_060294	NM_017824	4D	226394_at
chromosome 7 open reading frame 30	C7orf30	115416	NP_612455	NM_138446	4D	226385_s_at
chromosome 19 open reading frame 25	C19orf25	148223	NP_689695	NM_152482	4D	226378_s_at
TruB pseudouridine (psi) synthase homolog 1 (E. co	TRUBI	142940	NP_631908	NM_139169	4D	226339_at
6 hypothetical protein DKFZp762O076	DKFZp762O076	55529	NP_061180	NM_018710	4D	226338_at

640337 134957 937790 116236 689706 149076 689706 149076 937791 119587 001001936 84632 689807 161742 1116274 84971 653259 139818 689620 134266 689495 92370 149074 85403 376003 400924 006079 10928 060089 55599 775878 283635 065855 29993 065855 29993 065861 55140 000282 5230 006361 55140 006361 55140 00637 69503 257364 79133 006639 65268 694540 152518 0044178 57482 005716 10100 062824 25791 775889 283899 006058 1032	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540 XP_044178 NP_005716 NP_062824 NP_775889 NP_006058	4D NM_005725 4D NM_019850 4D NM_173618 4D NM_006067	227286_at 227363_s_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 149074 85403 EAF1 149074 85403 EAF1 149074 85403 EAF1 376003 400924 LOC400924 LOC400924 LOC400924 MCC40107 RNP 775878 283635 C14orf24 MGC855 29993 PACSINI MGC8979 254863 MGC40107 MGC92065 ELP3 MGC40107 BID MGC32065 WNK2 MGC32065 WNK2	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540 XP_044178 NP_062824 NP_062824 NP_775889		
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689495 139818 DOCK11 689495 13928 RAFEL2 689495 13928 RALBPI 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBPI 0063855 29993 RNP 775878 283635 C14orf24 065855 29993 PACSINI 689979 254863 MGC40107 695003 257364 MGC32065 077025 79133 C20orf7 694540 152518 HOZFP 0065716 10100 TSPAN2 005716	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540 XP_044178 NP_005716 NP_062824		
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24 06079 254863 MGC40107 000282 5230 PGK1 060561 55140 ELP3 0077025 79133 C20orf7 899062 119391 GSTO2 694503 65268 WNK2 0044178 5748	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540 XP_044178 NP_005716		
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 149078 10928 RALBP1 066089 55599 RNP 775878 283635 C14orf24 065855 29993 PACSINI 065861 55140 ELP3 0001187 637 BID 695003 257364 MGC32065 0077025 79133 C20orf7 694540 152518 HOZFP 044178 57482 KIAA1211	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540 XP_044178		227233_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 689820 139818 DOCK11 689495 92370 ACPL2 149074 85403 EAF1 149074 85403 EAF1 149074 85403 EAF1 106079 10928 RALBP1 060089 55599 RNP 065855 29993 PACSINI 689979 254863 MGC40107 060581 55140 ELP3 001187 637 BID 695003 257364 MGC32065 006639 65268 WNK2 694540 152518 HOZFP	NP_695003 NP_077025 NP_899062 NP_006639 NP_694540	4D XM_044178	227230_s_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAFI 376003 400924 LOC400924 006779 10928 RALBPI 060089 55599 RNP 775878 283635 C14orf24 060581 5230 PGK1 060581 55140 ELP3 060561 55140 ELP3 060563 257364 MGC32065 077025 79133 C20orf7 899062 119391 GSTO2 006639 65268 WNK2	NP_695003 NP_077025 NP_899062 NP_006639	4D NM_152995	227220_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 116274 84971 APG4D 689620 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006079 10928 RALBP1 065855 29993 PACSIN1 065855 29993 PACSIN1 000282 5230 PGK1 0007025 55140 ELP3 001187 637 BID 695003 257364 MGC32065 077025 79133 C20orf7 699062 119391 GSTO2	NP_695003 NP_077025 NP_899062	4D NM_006648	227217_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 149079 10928 RALBP1 0660089 55599 RNP 065855 29993 PACSINI 0658979 254863 MGC40107 060561 55140 ELP3 001187 637 BID 001187 637 BID 077025 79133 C20orf7	NP_695003 NP_077025	4D NM_183	227163_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24 065855 29993 PACSINI 060581 5230 PGK1 060561 55140 ELP3 001187 637 BID 695003 257364 MGC32065	NP_695003	4D NM_024	227160_s_at
640337 134957 RYESTRIB 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 116274 84971 APG4D 689620 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24 065855 29993 PACSINI 060581 55140 ELP3 0001187 637 BID		4D NM_153271	227151_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689495 92370 ACPL2 149074 85403 EAF1 149079 10928 RALBP1 0660089 55599 RNP 065855 29993 PACSINI 689979 254863 MGC40107 060561 55140 ELP3	NP_001187	4D NM_001196	227143_s_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 9377791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24 065855 29993 PACSINI 000282 5230 PGK1	NP_060561	4D NM_018091	227075_at
640337 134957 RYESTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 116274 84971 APG4D 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24 065855 29993 PACSIN1 689979 254863 MGC40107	NP_000282	4D NM_000291	227068_at
640337 134957 EWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689405 92370 ACPL2 149074 85403 EAF1 149079 10928 RALBP1 0660089 55599 RNP 065855 29993 PACSINI	NP_689979	4D NM_152766	227063_at -
640337 134957 FWISTRIA 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP 775878 283635 C14orf24	NP_065855	4D NM_020804	227053_at
640337 134957 TWISTRIB 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1 060089 55599 RNP	NP_775878	4D NM_173607	227029_at
640337 134957 TWISTRIB 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924 006779 10928 RALBP1	NP_060089	4D NM_017619	226975_at
640337 134957 FWISTRIB 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 901001936 84632 KIAA1914 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1 376003 400924 LOC400924	NP_006779	4D NM_006788	226957_x_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 937791 119587 CPXM2 937791 161742 SPREDI 689807 161742 SPREDI 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2 149074 85403 EAF1	XP_376003	4D XM_376003	226956_at
640337 134957 STXBP5 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11 689620 134266 GRPEL2 689495 92370 ACPL2	NP_149074	4D NM_033083	226952_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11	NP_689495	4D NM_152282	226925_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D 653259 139818 DOCK11	NP_689620	4D NM_152407	226881_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1 116274 84971 APG4D	NP_653259	4D NM_144658	226875_at
640337 134957 STXBPS 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914 689807 161742 SPRED1	NP_116274	4D NM_032885	226871_s_at '
640337 134957 STXBP5 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2 001001936 84632 KIAA1914	NP_689807	4D NM_152594	226837_at
640337 134957 STXBP5 937790 116236 LOC116236 689706 149076 FLJ25476 937791 119587 CPXM2	NP_001001936	4D NM_001001936	226829_at
640337 134957 STXBP5 937790 116236 LOC116236 689706 149076 FLJ25476	937791	4D NM_198	226824_at
640337 134957 STXBP5 937790 116236 LOC116236	689706	4D NM_15	226820_at
640337 134957 STXBP5	937790	4D NM_198	226796_at
OCTOOP WATER OCCUPANT	640337	4D NM_139	226794_at
001000006 201830 TWISTNIB	NP_001002926	4D NM_001002926	226784_at
NP_932068 154791 HSPC268 hypothetical protein HSPC268	NP_932068	4D NM_197964	226780_s_at
NP_112485 81889 FAHD1 fumarylacetoacetate hydrolase domain containing	NP_112485	4D NM_031208	226767_s_at

chromosome 20 open reading frame 149	C20orf149	79144	NP_077275	NM_024299	40	227994_x_at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	227942_s_at
exosome component 3	EXOSC3	51010	NP_057126	NM_016042	4D	227916_x_at
CDC42 effector protein (Rho GTPase binding) 5	CDC42EP5	148170	NP_659494	NM_145057	4D	227850_x_at
EPM2A (laforin) interacting protein 1	EPM2AIP1	9852	NP_055620	NM_014805	4D	227847_at
forminlike 3	FMNL3	91010	NP_783863	NM_175736	4D	227844_at
nudE nuclear distribution gene E homolog 1 (A. nid	NDEI	54820	NP_060138	NM_017668	4D	227843_at
zinc finger protein 605	ZNF605	90462	NP_899061	NM_183238	4D	227822_at
zinc finger protein 62 homolog (mouse)	ZFP62	92379	NP_689496	NM_152283	4D	227796_at
chromosome 6 open reading frame 157	C6orf157	90025	NP_944602	NM_198920	4D	227790_at
hypothetical protein FLJ90586	FLJ90586	135932	NP_699176	NM_153345	4D	227753_at
C3 and PZPlike, alpha2macroglobulin domain contain	CPAMD8	27151	NP_056507	NM_015692	4D	227721_at
suppressor of cytokine signaling 3	SOCS3	9021	NP_003946	NM_003955	4D	227697_at
similar to RIKEN cDNA 0610008P16 gene	MGC15937	92292	NP_542392	NM_080661	4D	227695_at
zinc finger protein 326	ZNF326	284695	NP_892021	NM_182976	4D	227680_at
Ku70binding protein 3	KUB3	91419	NP_150592	NM_033276	4D	227678_at
zinc finger protein 585A	ZNF585A	199704	NP_689868	NM_152655	4D	227674_at
doublecortin and CaM kinaselike 2	DCAMKL2	166614	NP_689832	NM_152619	4D	227666_at
Drosophil						
ELAV (embryonic lethal, abnormal vision,	ELAVL3	1995	NP_001411	NM_001420	4D	227612_at
epithelial stromal interaction 1 (breast)	EPSTII	94240	NP_001002264	NM_001002264	4D	227609_at
KIAA1627 protein	KIAA1627	57721	NP_066012	NM_020961	4D	227601_at
hypothetical protein FLJ12949	FLJ12949	65095	NP_075384	NM_023008	4D	227587_at
hypothetical protein MGC42630	MGC42630	286301	NP_787119	NM_175923	4D	227563_at
PHD finger protein 20like 1	PHF20L1	51105	NP_057102	NM_016018	4D	227523_s_at
Fbox protein 33	FBXO33	254170	NP_976046	NM_203301	4D	227521_at
forkhead box Q1	FOXQ1	94234	NP_150285	NM_033260	4D	227475_at
HEC1 domain and ankyrin repeat containing, E3 ubiq	HACEI	3/331	NF_063822	NM_020//1	45	22/4/1_at
hypothetical protein LOC197322	LOC197322	197322	NP_777577	NM_174917	#D	227464_at
EPH receptor A4	EPHA4	2043	NP_004429	NM_004438	4D	227449_at
hypothetical protein FLJ38991	FLJ38991	285521	NP_776188	NM_173827	4D	227442_at
hypothetical protein MGC14595	MGC14595	84294	NP_115710	NM_032334	4D	227402_s_at

9899 at JD NM 138499 N.P. 617508 LP7034 LPWRP2 PWRWP domain containing 2 98097 x at 44D NM 013586 NP 695411 308.44 ZRMD1 zinc rebon domain containing 2 9807 x at 44D NM 019922 NP 696422 55337 FL20788 EL207038 protein 8805 x at 44D NM 01019790 NP 696422 55337 FL207788 FL207038 protein 8805 x at 44D NM 01019790 NP 975273 388888 LOC288888 Inchronsome 90 pone reading frame 105 8805 x at 44D NM 010744 NP 997527 388888 LOC288888 Inchronsome 90 pone reading frame 105 8807 x at 44D NM 120361 NP 975827 388144 MCC3907 Bell CLL/symphoma 9like 8809 x at 44D NM 120363 NP 975827 388149 BCLV Bell CLL/symphoma 9like 8810 at 44D NM 1203231 NP 904963 162972 ZNF580 zinc finger protein 550 8810 at 4D NM 138983 NP 904943 162	chromosome 21 open reading frame 45	C21orf45	54069	NP_061817	NM_018944	4D	228597_at
at 4D NM_138499 NP_612508 17034 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 x at 4D NM_014596 NP_055411 30834 ZNRD1 z at 4D NM_0172864 NP_060603 128414 C20ort58 z at 4D NM_0170792 NP_060602 5037 FL20758 z at 4D NM_010010790 NP_001001790 401505 C90rf105 z at 4D NM_0182557 NP_872363 283149 BCL9L z at 4D NM_182527 NP_907227 38886 LOC388886 z at 4D NM_193337 NP_6935369 374395 LOC374395 x at 4D NM_017341 NP_903536 NP_20185 CACNA2D4 x at 4D NM_017341 NP_90353 NP_20365 LOC374395 x at 4D NM_017341 NP_904963 16248 PL20280 x at 4D NM_02323	FLJ20758 protein	FLJ20758	55037	NP_060422	NM_017952	4D	228590_at
at 4D NM_138499 NP_612508 17034 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_01952 NP_060603 128414 C20orf58 at 4D NM_01952 NP_060603 28314 C20orf58 at 4D NM_01952 NP_00101790 401505 C9orf105 at 4D NM_01962 NP_00101790 40150 BCL9L at 4D NM_01932 NP_58952 93589 CACNA2D4 x at 4D NM_102324 NP_978952 93589 CACNA2D4 x at 4D NM_103288 NP_115661 84245 MGC3207 at 4D NM_103289 NP_115661 84248 CACNA2D4 x at 4D NM_133983 NP_060211 54876 FLJ20280 x at 4D NM_032896 NP_115644	CSRP2 binding protein	CSRP2BP	57325	NP_065397	NM_020536	4D	228544_s_at
at 4D NM_138499 NP_612508 17034 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_0164596 NP_050603 128414 CZ0orf58 at 4D NM_01010790 NP_060603 128414 CZ0orf58 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_182557 NP_15661 84248 MCG3207 at 4D NM_172364 NP_2758952 93589 CACNA2D4 xat 4D NM_172364 NP_285369 374395 LOC34395 at 4D NM_172364 NP_285369 374395 LOC34395 xat 4D NM_0132883	CSRP2 binding protein	CSRP2BP	57325	NP_065397	NM_020536	4D	228543_at
at 4D NM_138499 NP_612508 17034 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_0104590 NP_050603 128414 C20orf58 at 4D NM_001001790 NP_0000279 401505 C9orf105 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_001001790 NP_001001790 40150 C9orf105 at 4D NM_001001790 NP_001001790 40150 C9orf105 at 4D NM_182537 NP_001001790 40150 CACNA2D4 x_at 4D NM_172364 NP_058569 374395 LOC374395 at 4D NM_172364 NP_058569 374395 LOC374395 at 4D NM_173231 NP_060287 374395 LOC374395 at 4D NM_0323	sterile alpha motif domain containing 9	SAMD9	54809	NP_060124	NM_017654	4D	228531_at
at 4D NM_138499 NP_612508 17034 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_017952 NP_690603 128414 C20orf38 at 4D NM_017952 NP_600422 55037 FL20758 at 4D NM_017952 NP_97527 38886 COorf105 at 4D NM_0170644 NP_97527 38886 MGC3207 at 4D NM_0172364 NP_975892 93889 CACNA2D4 x at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_032337 NP_953369 374395 LOC374395 at 4D NM_0173241 NP_904963 162972 ZNF550 at 4D NM_023883 NP_604963 162972 ZNF550 x at 4D NM_032898 NP_16249	chromosome 10 open reading frame 86	C10orf86	54780	NP_060085	NM_017615	4D	228506_at
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_050603 128414 C20orf38 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_01707644 NP_97527 38886 LOC38886 at 4D NM_182557 NP_872363 283149 BCLD1 at 4D NM_032285 NP_115661 84245 MCC3207 at 4D NM_032337 NP_953369 374395 LOC374395 at 4D NM_0173264 NP_90327 1390 CREM at 4D NM_0173289 NP_904963 162972 ZNF550 at 4D NM_032898 NP_115684	THAP domain containing 8	THAP8	199745			4D	228500_at
att 4D NM_138499 NP_612508 170394 PWWP2 x_att 4D NM_014596 NP_055411 30834 ZNRD1 att 4D NM_014596 NP_050421 30834 ZNRD1 att 4D NM_017952 NP_060422 55037 FLJ20758 att 4D NM_017952 NP_001001790 401505 C9orf105 att 4D NM_017934 NP_997527 38886 LOC388886 att 4D NM_0182557 NP_872363 283149 BCL9L att 4D NM_022285 NP_115661 84245 MGC3207 at 4D NM_172364 NP_997527 38886 LOC378886 y at 4D NM_19337 NP_694963 18245 MGC3207 at 4D NM_013288 NP_694963 162972 ZNF550 at 4D NM_013289 NP_604963 162972 ZNF550 at 4D NM_023293 NP_60496	6phosphofructo2kinase/fructose2,6biphosphatase 4	PFKFB4	5210	NP_004558	NM_004567	4D	228499_at
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 at 4D NM_017952 NP_060422 55037 FLJ20758 at 4D NM_017952 NP_01001790 401505 C9orf105 at 4D NM_01707644 NP_97527 38886 LOC388886 at 4D NM_027644 NP_97527 38886 LOC388886 at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_694963 29389 CACNA2D4 x at 4D NM_133898 NP_001872 3389 CACNA2D4 x at 4D NM_133898 NP_694963 162972 ZNF550 at 4D NM_032380 NP_16274035 4629 MYH11 x at 4D NM_032398 NP_2674035					NM_018675	4D	
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_060422 128414 ZNRD1 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_017952 NP_060422 55037 FLJ20758 at 4D NM_017952 NP_060422 55037 FLJ20758 at 4D NM_0101790 NP_060422 55037 FLJ20758 at 4D NM_0101790 NP_00101790 401505 C9orf105 at 4D NM_182557 NP_00101790 401505 C9orf105 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_1932557 NP_115661 84245 MGC3207 at 4D NM_193337 NP_953369 374395 LOC374395 at 4D NM_193231 NP_90	alphakinase 2	ALPK2	115701		NM_052947	4D	228367_at
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 5037 FL20758 at 4D NM_017952 NP_01001790 401505 C9orf105 at 4D NM_017952 NP_97527 38886 LOC38886 at 4D NM_0170644 NP_975263 283149 BCL9L at 4D NM_0232285 NP_115661 84245 MCC3207 at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_193231 NP_955369 374395 LOC374395 at 4D NM_1383231 NP_901872 1390 CREM Ap 4D NM_13838 NP_604963 <td>hypothetical protein LOC91942</td> <td>LOC91942</td> <td>91942</td> <td>1 .</td> <td></td> <td>4D</td> <td></td>	hypothetical protein LOC91942	LOC91942	91942	1 .		4D	
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_060422 55037 FL20758 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_017952 NP_060422 55037 FL20758 at 4D NM_0170644 NP_975273 38886 LOC38886 at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_193236 NP_958369 374395 LOC374395 at 4D NM_153231 NP_0694963 162972 ZNF550 at 4D NM_0138983 NP_0694963 162972 ZNF550 at 4D NM_032308 NP_115684 84268 RIP at 4D NM_032380 NP_116279	hypothetical protein MGC16044	MGC16044	91523	11		4D	228298_at
at 4D NM_138499 NP_612508 170394 PWWP2 x at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_014596 NP_060422 55037 FL20758 s_at 4D NM_017952 NP_060422 55037 FL20758 s_at 4D NM_017952 NP_001001790 401505 C9orf105 at 4D NM_0182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_193337 NP_953369 374395 LOC374395 at 4D NM_153231 NP_953369 374395 LOC374395 at 4D NM_0138983 NP_964963 162972 ZNF550 at 4D NM_0138983 NP_964963 162972 ZNF550 at 4D NM_032890 NP_964963 162972 ZNF550 at 4D NM_032893 NP_964	novel 58.3 KDA protein	LOC91614	91614		NM_139160	4D	228293_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf58 s_at 4D NM_017952 NP_060422 55037 FLJ20758 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_007644 NP_97527 38886 LOC388886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_172364 NP_9578952 9389 CACNA2D4 x_at 4D NM_199337 NP_95369 374395 LOC374395 at 4D NM_0153231 NP_9694963 162972 ZNF550 at 4D NM_017741 NP_060211 54876 FLJ20280 at 4D NM_032890 NP_115684 84268 RIP s_at 4D NM_03289 NP_15	tudor domain containing 9	TDRD9	122402	NP_694591	NM_153046	4D	228285_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_060422 55037 FL20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_001001790 NP_97527 38886 LOC388886 at 4D NM_182557 NP_972363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_953952 93589 CACNA2D4 x_at 4D NM_199337 NP_95369 374395 LOC374395 at 4D NM_001881 NP_901872 1390 CREM at 4D NM_0153231 NP_694963 162972 ZNF550 at 4D NM_0138983 NP_60211 54876 FL20280 at 4D NM_032890 NP_11668	l'aminocyclopropane l'carboxylate synthase	PHACS	84680	NP_115981	NM_032592	4D	228264_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_017952 NP_97527 38886 LOC38886 at 4D NM_032285 NP_15661 84245 MGC3207 at 4D NM_193257 NP_58952 93589 CACNA2D4 x_at 4D NM_193337 NP_95369 374395 LOC374395 at 4D NM_193231 NP_95369 374395 LOC374395 at 4D NM_153231 NP_694963 162972 ZNF550 at 4D NM_017741 NP_060211 54876 FLJ20280 at 4D NM_032898 NP_115684 84268 RIP at 4D NM_032389 NP_115684	proline argininerich end leucinerich repeat protei	PRELP	5549	NP_002716	NM_002725	4D	228224_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_01752864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_0182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_172364 NP_758952 93589 CACNA2D4 x_at 4D NM_199337 NP_95369 374395 LOC374395 at 4D NM_001881 NP_001872 1390 CREM at 4D NM_0153231 NP_694963 162972 ZNF550 at 4D NM_0138983 NP_600211 54876 FLJ20280 s_at 4D NM_032308 NP	hypothetical LOC389362	LOC389362	389362	XP_374159	XM_374159	4D	228217_s_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_017964 NP_997527 38886 LOC388886 at 4D NM_032285 NP_872363 283149 BCL9L at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_192364 NP_758952 93589 CACNA2D4 x_at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_153231 NP_955369 374395 LOC374395 at 4D NM_017741 NP_001872 1390 CREM at 4D NM_0138983 NP_00211 54876 FLJ20280 at 4D NM_0138983 NP_004035 </td <td>dispatched homolog 1 (Drosophila)</td> <td>DISP1</td> <td>84976</td> <td>NP_116279</td> <td>NM_032890</td> <td>4D</td> <td>228184_at</td>	dispatched homolog 1 (Drosophila)	DISP1	84976	NP_116279	NM_032890	4D	228184_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf38 at 4D NM_017952 NP_060422 55037 FLJ20758 at 4D NM_001001790 NP_997527 38886 LOC388866 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_0183231 NP_964963 162972 ZNF550 at 4D NM_022844 NP_060211 54876 FLJ20280 at 4D NM_022844 NP_074035 4629 MYH11 at 4D NM_138983 NP_620450 116448 OLIG1	RPA interacting protein	RIP	84268		NM_032308	4D	
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf38 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_0207644 NP_997527 38886 LOC388886 at 4D NM_032285 NP_115661 84245 MCC3207 at 4D NM_172364 NP_758952 93589 CACNA2D4 x_at 4D NM_199337 NP_955369 374395 LOC374395 x_at 4D NM_001881 NP_955369 374395 LOC374395 at 4D NM_001881 NP_001872 1390 CREM at 4D NM_017341 NP_060211 54876 FLJ20280 at 4D NM_017741 NP_074035 4629 MYH11	oligodendrocyte transcription factor 1	OLIG1	116448	NP_620450	NM_138983	4D	228170_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_207644 NP_997527 38886 LOC38886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_95369 374395 LOC374395 x_at 4D NM_199331 NP_901872 1390 CREM at 4D NM_001881 NP_001872 1390 CREM at 4D NM_017741 NP_060211 54876 FLJ20280	myosin, heavy polypeptide 11, smooth muscle	MYHII	4629	E 1	NM_022844	4D	228134_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf58 at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_207644 NP_997527 388886 LOC388886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_172364 NP_955369 374395 CACNA2D4 x_at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_1091881 NP_001872 1390 CREM At 4D NM_153231 NP_604963 162972 ZNF550	hypothetical protein FLJ20280	FLJ20280	54876	11	NM_017741	4D	228106_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 s_at 4D NM_207644 NP_997527 38886 LOC38886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_955369 374395 LOC374395 at 4D NM_199337 NP_955369 374395 LOC374395	zinc finger protein 550	ZNF550	162972		11 1	4D	228099_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf58 at 4D NM_001001790 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_997527 38886 LOC38886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_199337 NP_955369 374395 LOC374395	cAMP responsive element modulator	CREM	1390		NM_001881	4D	228092_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_0017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_207644 NP_997527 388886 LOC388886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207 at 4D NM_172364 NP_758952 93589 CACNA2D4	similar to RIKEN cDNA 1810059G22	LOC374395	374395	NP_955369	NM_199337	4D	228089_x_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_017952 NP_690603 128414 C20orf58 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 s_at 4D NM_207644 NP_997527 388886 LOC388886 at 4D NM_182557 NP_872363 283149 BCL9L at 4D NM_032285 NP_115661 84245 MGC3207	calcium channel, voltagedependent, alpha 2/delta s	CACNA2D4	93589		NM_172364	4D	228083_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_207644 NP_997527 388886 LOC388886 at 4D NM_182557 NP_872363 283149 BCL9L	hypothetical protein MGC3207	MGC3207	84245		NM_032285	4D	228077_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105 at 4D NM_207644 NP_997527 388886 LOC388886	Bcell CLL/lymphoma 9like	BCL9L	283149	NP_872363	NM_182557	4D	228065_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758 s_at 4D NM_001001790 NP_001001790 401505 C9orf105	similar to hypothetical protein LOC192734	LOC388886	388886	3	NM_207644	4D	228064_at
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58 at 4D NM_017952 NP_060422 55037 FLJ20758	chromosome 9 open reading frame 105	C9orf105	401505	,	NM_001001790	4D	
at 4D NM_138499 NP_612508 170394 PWWP2 x_at 4D NM_014596 NP_055411 30834 ZNRD1 s_at 4D NM_152864 NP_690603 128414 C20orf58	FLJ20758 protein	FLJ20758	55037		NM_017952	4D	228020_at
at 4D NM_138499 NP_612508 170394 PWWP2 _x_at 4D NM_014596 NP_055411 30834 ZNRD1	chromosome 20 open reading frame 58	C20orf58	128414		1 1	4D	1
at 4D NM_138499 NP_612508 170394 PWWP2	zinc ribbon domain containing, 1	ZNRDI	30834			4D	1
	PWWP domain containing 2	PWWP2	170394	1 1		4D	227999_at

Tcell activation GTPase activating protein	TAGAP	117289	NP_473455	NM_054114	40	229/23_at
cleavage stimulation factor, 3' preRNA, subunit 3,	CSTF3	1479	NP_001317	NM_001326		229665_at
peroxisomal lon protease	LONP	83752	NP_113678	NM_031490		229663_at
chromosome 18 open reading frame 51	C18orf51	125704	XP_058931	XM_058931	4D	229645_at
iroquois homeobox protein 3	IRX3	79191	NP_077312	NM_024336	4D	229638_at
hypothetical protein FLJ32752	FLJ32752	144132	NP_653267	NM_144666		
hypothetical protein FLJ39441	FLJ39441	144108	NP_919261	NM_194285	4D	229594_at
Tbox 3 (ulnar mammary syndrome)	ТВХЗ	6926	NP_005987	NM_005996	4D	229576_s_at
peroxisomal acylCoA thioesterase 2B	PTE2B	122970	NP_689544	NM_152331	4D	229534_at
fragile X mental retardation, autosomal homolog 1	FXRI	8087	NP_005078	NM_005087	4D	229519_at
family with sequence similarity 46, member B	FAM46B	115572	NP_443175	NM_052943	4D	229518_at
cytochrome c oxidase subunit Va	COX5A	9377	NP_004246	NM_004255	4D	229426_at
golgi transport 1 homolog A (S. cerevisiae)	GOLTIA	127845	NP_940849	NM_198447	4D	229372_at
immunoglobulin superfamily, member 4C	IGSF4C	199731	NP_660339	NM_145296	4D	229335_at
hypothetical protein MGC2749	MGC2749	79036	NP_076974	NM_024069	4D	229316_at
junctophilin 3	JPH3	57338	NP_065706	NM_020655	4D	229294_at
ribonuclease L (2',5'oligoisoadenylate synthetased	RNASEL	6041	NP_066956	NM_021133	4D	229285_at
collagen, type XI, alpha 1	COLITAI	1301	NP_001845	NM_001854	4D	229271_x_at
LOC440667	LOC440667	440667	XP_498810	XM_498810	4D	229264_at
lactate dehydrogenase D	LDHD	197257	NP_705690	NM_153486	4D	229241_at
KIAA0251 protein	KIAA0251	23042	NP_055842	NM_015027	4D	229219_s_at
solute carrier family 35, member F3	SLC35F3	148641	NP_775779	NM_173508	4D	229065_at
hypothetical protein MGC31967	MGC31967	203260	NP_777583	NM_174923	4D	229063_s_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT17	200035	1	XM_496395	40	1 1
zinc finger protein 449	ZNF449	203523	NP_689908	NM_152695	4D	228968_at
hypothetical protein FLJ35954	FLJ35954	166968	NP_689835	NM_152622	4D	228961_at
S100 calcium binding protein A6 (calcyclin)	S100A6	6277	NP_055439	NM_014624	4D	228923_at
prematurely terminated mRNA decay factorlike	LOC91431	91431	NP_619643	NM_138698	4D	228859_at
hypothetical protein LOC196463	LOC196463	196463	NP_775813	NM_173542	4D	228845_at
solute carrier family 13 (sodiumdependent citrate	SLC13A5	284111	NP_808218	NM_177550	4D	228844_at
KIAA1571 protein	KIAA1571	57683	XP_371590	XM_371590	4D	228749_at
zinc finger, CCHC domain containing 12	ZCCHC12	170261	NP_776159	NM_173798	4D	228715_at
procollagenproline, 2oxoglutarate 4dioxygenase (pr	P4HA3	283208	NP_878907	NM_182904	4D	228703_at

hypothetical protein BC014608	LOC128153	128153	NP_620151	NM_138796	4D	230763_at
hypothetical protein MGC33414	MGC33414	257101	NP_775845	NM_173574	4D	230756_at
hypothetical protein LOC146174	LOC146174	146174	NP_775772	NM_173501	4D	230721_at
kinesin family member 9	KIF9	64147	NP_071737	NM_022342	4D	230701_x_at
hypothetical STGC3	STGC3	474171	NP_001006609	NM_001006608	4D	230587_at
hypothetical LOC388480	LOC388480	388480	XP_373779	XM_373779	4D	230574_at
cAMP responsive element modulator	CREM	1390	NP_874391	NM_182722	4D	230511_at
CD5 antigen (p5662)	CD5	921	NP_055022	NM_014207	4D	230489_at
similar to RIKEN cDNA 6030419C18 gene	LOC388135	388135	XP_370873	XM_370873	4D	230475_at
hypothetical gene supported by AK124342	LOC400999	400999	XP_379123	XM_379123	4D	230452_at
similar to hypothetical protein 9630041N07	LOC345462	345462	XP_293828	XM_293828	4D	230421_at
inositol polyphosphate5phosphatase F	INPP5F	22876	NP_938145	NM_198331	4D	230363_s_at
inositol polyphosphate5phosphatase F	INPPSF	22876	NP_938145	NM_198331	4D	230362_at
transmembrane protein 45B	TMEM45B	120224	NP_620143	NM_138788	4D	230323_s_at
chromosome 9 open reading frame 10 opposite strand	C9orf10OS	158293	NP_942138	NM_198841	4D	230308_at
RAB7B, member RAS oncogene family	RAB7B	338382	NP_796377	NM_177403	4D	230266_at
ST8 alphaNacetylneuraminide alpha2,8sialyltransfer	ST8SIA4	7903	NP_005659	NM_005668	4D	230261_at
placentaspecific 9	PLAC9	219348	XP_166090	XM_166090	4D	230246_at
ret finger protein 2	RFP2	10206		NM_005798	4D	230192_at
family with sequence similarity 14, member B	FAM14B	122509	NP_660292	NM_145249	4D	230172_at
similar to Histone H3.3	LOC392533	392533	XP_373368	XM_373368	4D	230144_at
solute carrier family 30 (zinc transporter), membe	SLC30A2	7780	NP_115902	NM_032513	4D	230084_at
BTB (POZ) domain containing 14A	BTBD14A	138151	NP_653254	NM_144653	4D	230050_at
mitochondrial ribosomal protein L43	MRPL43	84545	NP_789763	NM_176793	4D	230026_at
hypothetical protein FLJ21103	FLJ21103	79607	NP_078832	NM_024556	4D	230009_at
tigger transposable element derived 2	TIGD2	166815	NP_663761	NM_145715	4D	229983_at
Ellis van Creveld syndrome 2 (limbin)	EVC2	132884	NP_667338	NM_147127	4D	229974_at
FLJ32363 protein	FLJ32363	375444	NP_940968	NM_198566	4D	229886_at
Kruppellike factor 12	KLF12	11278	NP_009180	NM_007249	4D	229881_at
ventricular zone expressed PH domain homolog 1 (ze	VEPH1	79674	NP_078897	NM_024621	4D	229/59_s_at
hypothetical LOC145853	LOC145853	145853	XP_096885	XM_096885	4D	229742_at
similar to RIKEN cDNA 1700027J05 gene	MGC33692	414918	NP_001001794	NM_001001794	4D	229739_s_at

RNA (guanine9) methyltransferase domain containing	RG9MTD2	93587	NP_689505	NM_152292	4D	231877_at
kinesin family member 21A	KIF21A	55605	NP_060111	NM_017641	4D	231875_at
inhibitor of growth family, member 3	ING3	54556	NP_938008	NM_198267	4D	231863_at
tubulin, delta 1	TUBD1	51174	NP_057345	NM_016261	4D	231853_at
alanyltRNA synthetase like	AARSL	57505	NP_065796	NM_020745	4D	231845_at
KIAA1272 protein	KIAA1272	57186	XP_046600	XM_046600	4D	231826_at
FLJ10378 protein	FLJ10378	55132	NP_060548	NM_018078	4D	231824_at
noggin	NOG	9241	NP_005441	NM_005450	4D	231798_at
distalless homeo box 3	DLX3	1747	NP_005211	NM_005220	4D	231778_at
member	INFRSFIOA	8/9/	NP_003835	NM_003844	4D	231775_at
upstream transcription factor 1	USFI	7391	NP_009053	NM_007122	4D	231768_at
zona pellucida glycoprotein 4	ZP4	57829	NP_067009	NM_021186	4D	231756_at
winglesstype MMTV integration site family, member	WNT3	7473	NP_110380	NM_030753	4D	231743_at
tissue inhibitor of metalloproteinase 2	TIMP2	7077	NP_003246	NM_003255	4D	231579_s_at
guanylate binding protein 1, interferoninducible,	GBP1	2633	NP_002044	NM_002053	4D	231578_at
chromosome 9 open reading frame 138	C9orf138	158297	NP_714918	NM_153707	4D	231570_at
hypothetical protein MGC35194	MGC35194	93190	NP_689503	NM_152290	4D	231549_at
chromosome 11 open reading frame 1	Cllorfi	64776	NP_073598	NM_022761	4D	231530_s_at
chromosome 20 open reading frame 144	C20orf144	128864	NP_543015	NM_080825	4D	231522_at
gametogenetin	GGN	199720	NP_689870	NM_152657	4D	231420_at
LOC122258	LOC122258	122258	NP_660291	NM_145248	4D	231405_at
carbonic anhydrase XIII	CA13	377677	NP_940986	NM_198584	4D	231270_at
similar to RIKEN cDNA 1110018M03	LOC387758	387758	NP_976249	NM_203371	4D	231001_at
Bcell novel protein 1	BCNP1	199786	NP_775815	NM_173544	4D	230983_at
HCG3 gene	HCG3	414061	NP_001001394	NM_001001394	4D	230953_at
cerevis					Ī	## 0000 F
CDC14 cell division cycle 14 homolog B (S.	CDC14B	8555	NP 201589	NM 033332	4D	230887 at
BCL2associated athanogene 2	BAG2	9532	NP_004273	NM 004282	4D	230879 at
cystatin 9like (mouse)	CST9L	128821	NP_542177	NM_080610	4D	230829 at
54 SVAP1 protein	IMAGE3451454	116841	NP_444280	NM_053052	4D	230810 at
FLJ37099 protein	FLJ37099	163259	NP_940861	NM_198459	4D	230769 at

hypothetical protein FLJ23322	FLJ23322	80020	NP_079231	NM_024955	4D	233250_x_at
general transcription factor IIIC, polypeptide 5,	GTF3C5	9328	NP_036219	NM_012087	4D	233173_x_at
UDPNacetylalphaDgalactosamine:polypeptide Nacetylg	GALNTL5	168391	NP_660335	NM_145292	4D	233150_at
chromosome 20 open reading frame 142	C20orf142	128486	XP_371399	XM_371399	4D	233061_at
chromosome X open reading frame 44	CXorf44	90736	NP_612371	NM_138362	4D	232990_at
hypothetical protein FLJ20444	FLJ20444	403323	NP_982274	NM_203449	4D	232897_at
chromosome 20 open reading frame 166	C20orf166	128826	NP_848558	NM_178463	4D	232892_at
dedicator of cytokinesis 8	DOCK8	81704	NP_982272	NM_203447	4D	232843_s_at
PWWP domain containing 1	PWWPI	154150	NP_612641	NM_138574	4D	232680_at
hypothetical gene supported by AK022396; AK097927	LOC400946	400946	XP_379079	XM_379079	4D	232664_at
APG16 autophagy 16like (S. cerevisiae)	APG16L	55054	NP_060444	NM_017974	4D	232612_s_at
WAP fourdisulfide core domain 3	WFDC3	140686	NP_542181	NM_080614	4D	232602_at
hypothetical protein MGC27466	MGC27466	127396	NP_689586	NM_152373	4D	232563_at
leucine rich repeat and fibronectin type III domai	LRFN1	57622	XP_290842	XM_290842	4D	232486_at
chromosome 10 open reading frame 71	C10orf71	118461	NP_955629	NM_199459	4D	232456_at
hypothetical protein FLJ21439	FLJ21439	80208	NP_079413	NM_025137	4D	232403_at
PDZ and LIM domain 2 (mystique)	PDLIM2	64236	NP_789847	NM_176871	4D	232302_at
piggyBac transposable element derived 3	PGBD3	267004	NP_736609	NM_170753	4D	232287_at
solute carrier family 6, member 15	SLC6A15	55117	NP_060527	NM_018057	4D	232263_at
amyotrophic lateral sclerosis 4	ALS4	23064	NP_055861	NM_015046	4D	232229_at
KIAA1465 protein	KIAA1465	57611	NP_065902	NM_020851	4D	232208_at
palmdelphin	PALMD	54873	NP_060204	NM_017734	4D	232187_at
serine active site containing 1	SERAC1	84947	NP_116250	NM_032861	4D	232183_at
hypothetical protein FLJ14129	FLJ14129	79943	NP_112157	NM_030895	4D	232162_at
subun			1	ı		
mediator of RNA polymerase II transcription,	MED12L	116931	NP_443728	NM 053002	4D	232114 at
similar to hypothetical protein Y97E10AL.1	DKFZp761P211	375057	NP_001003665	NM_001003665	4D	232104 at
chromosome X open reading frame 23	CXorf23	256643	NP_938020	NM_198279	4D	232087_at
Nedd4 binding protein 2	N4BP2	55728	NP_060647	NM_018177	4D	231996_at
chromosome 20 open reading frame 160	C20orf160	140706	NP_542192	NM_080625	4D	231991_at
similar to HSPC323	LOC284422	284422	XP_209196	XM_209196	4D	231982_at
PHD finger protein 20like 1	PHF20L1	51105	NP_940915	NM_198513	4D	231967_at
zinc finger protein 529	ZNF529	57711	NP_066002	NM_020951	4D	231940_at
Some assumption of the control of th						

deoxynucleotidyltransferase, terminal, interacting	DNTTIPI	116092	NP_443183	NM_052951	4D	234942_s_at
ribosomal protein L7a	RPL7A	6130	NP_000963	NM_000972	4D	234873_x_at
transient receptor potential cation channel, subfa	TRPM6	140803	NP_060132	NM_017662	4D	234864_s_at
G2 protein	G2	25758	XP_039515	XM_039515	4D	234784_at
KIAA1596	KIAA1596	57697	XP_048128	XM_048128	4D	234733_s_at
poly (ADPribose) polymerase family, member 6	PARP6	56965	NP_064598	NM_020213	4D	234710_s_at
similar to bA182L21.1 (novel protein similar to hy	LOC399788	399788	XP_374817	XM_374817	4D	234695_x_at
olfactory receptor, family 52, subfamily D, member	OR52D1	390066	NP_001005163	NM_001005163	4D	234617_at
likely ortholog of mouse NOBOX	NOBOX	135935	XP_069612	XM_069612	4D	234519_at
sperm associated antigen 4like	SPAG4L	140732	NP_542406	NM_080675	4D	234504_at
cytotoxic Tlymphocyteassociated protein 4	CTLA4	1493	NP_005205	NM_005214	4D	234362_s_at
zinc finger protein 326	ZNF326	284695	NP_892020	NM_182975	4D	234093_at
serine/threonine kinase 36 (fused homolog, Drosoph	STK36	27148	NP_056505	NM_015690	4D	234005_x_at
pleckstrin homology domain containing, family G (w	PLEKHG2	64857	NP_073746	NM_022835	4D	233986_s_at
espin	ESPN	83715	NP_113663	NM_031475	4D	233979_s_at
CGI09 protein	CG109	51605	NP_057023	NM_015939	4D	233970_s_at
cystatin 11	CST11	140880	NP_570612	NM_130794	4D	233968_at
CXYorf1related protein	FLJ25222	374666	NP_954632	NM_199163	4D	233929_x_at
SLITROBO Rho GTPase activating protein 1	SRGAPI	57522	NP_065813	NM_020762	4D	233888_s_at
chromosome 20 open reading frame 43	C20orf43	51507	NP_057491	NM_016407	4D	233842_x_at
ATbinding transcription factor 1	ATBF1	463	NP_008816	NM_006885	4D	233751_at
keratin associated protein 94	KRTAP94	85280	NP_149461	NM_033191	4D	233640_x_at
signalinduced proliferationassociated 1 like 2	SIPA1L2	57568	NP_065859	NM_020808	4D	233587_s_at
chromosome 20 open reading frame 149	C20orf149	79144	NP_077275	NM_024299	4D	233571_x_at
cofilin 2 (muscle)	CFL2	1073	NP_068733	NM_021914	4D	233496_s_at
hypothetical protein LOC96597	LOC96597	96597	XP_378655	XM_378655	4D	233483_at
zinc finger protein 226	ZNF226	7769	NP_057528	NM_016444	4D	233461_x_at
polymerase (DNA directed) nu	POLN	353497	NP_861524	NM_181808	4D	233454_at
DEP domain containing 2	DEPDC2	80243	NP_079146	NM_024870	4D	233408_at
pericentrin 2 (kendrin)	PCNT2	5116	NP_006022	NM_006031	4D	233387_s_at
polymerase (RNA) I polypeptide B, 128kDa	POLR1B	84172	NP_061887	NM_019014	4D	233341_s_at
churchill domain containing 1	CHURC1	91612	NP_660148	NM_145165	4D	233268_s_at

vesicle transport through interaction with	VTIIA	143187	NP_660207	NM_145206	40	235923_at
guanylate binding protein 4	GBP4	115361	NP_443173		4D	235574_at
hypothetical protein MGC40214	MGC40214	137682	NP_689629	NM_152416	4D	235509_at
ankyrin repeat and SOCS boxcontaining 5	ASB5	140458	NP_543150	NM_080874	4D	235503_at
fucosyltransferase 10 (alpha (1,3) fucosyltransfer	FUT10	84750	NP_116053	NM_032664	4D	235472_at
TruB pseudouridine (psi) synthase homolog 1 (E. co	TRUB1	142940	NP_631908	NM_139169	4D	235447_at
shugoshinlike 2 (S. pombe)	SGOL2	151246	NP_689737	NM_152524	4D	235425_at
similar to RP2 protein, testosteroneregulated	LOC390916	390916	XP_372723	XM_372723	4D	235384_at
ELLP3030	UNQ3030	375387	NP_940967	NM_198565	4D	235359_at
UBX domain containing 4	UBXD4	165324	NP_859064	NM_181713	4D	235327_x_at
RNA (guanine9) methyltransferase domain containing	RG9MTD3	158234	NP_659401	NM_144964	4D	235303_at
eukaryotic translation initiation factor 5A2	EIF5A2	56648	NP_065123	NM_020390	4D	235296_at
bone morphogenetic protein 8b (osteogenic protein	BMP8B	656	NP_001711	NM_001720	4D	235275_at
DCP2 decapping enzyme homolog (S. cerevisiae)	DCP2	167227	NP_689837	NM_152624	4D	235258_at
folliculin	FLCN	201163	NP_653207	NM_144606	4D	235250_at
hypothetical protein FLJ90709	FLJ90709	153129	NP_775785	NM_173514	4D	235241_at
similar to hepatocellular carcinomaassociated anti	LOC151194	151194	NP_660323	NM_145280	4D	235177_at
TAF3 RNA polymerase II, TATA box binding protein (TAF3	83860	XP_291729	XM_291729	4D	235119_at
hook homolog 3 (Drosophila)	ноок3	84376	NP_115786	NM_032410	4D	235114_x_at
HRASlike suppressor 3	HRASLS3	111145	NP_009000	NM_007069	4D	235110_at
Fbox and leucinerich repeat protein 20	FBXL20	84961	NP_116264	NM_032875	4D	235089_at
nuclear domain 10 protein	NDP52	10241	NP_005822	NM_005831	4D	235076_at
transmembrane protein 41A	TMEM41A	90407	NP_542383	NM_080652	4D	235037_at
similar to Zinc finger protein 58 (Zfp58) (Zinc fi	LOC402524	402524	XP_379850	XM_379850	4D	235027_at
TAF4b RNA polymerase II, TATA box binding protein	l'AF4B	68/3	XP_290809	XM_290809	40	235020_at
BardetBiedl syndrome 7	BBS7	55212	NP_789794	NM_176824	4D	235007_at
valosincontaining protein (p97)/p47 complexinterac	VCIP135	80124	NP_079330	NM_025054	4D	234988_at
hypothetical protein LOC143458	LOC143458	143458	NP_777562	NM_174902	4D	234985_at
similar to Zincalpha2glycoprotein precursor (Znalp	LOC401393	401393	XP_376658	XM_376658	4D	234955_at

TEX13A testis expressed sequence 13A 143 MGC50372 hypothetical protein MGC50372 468 C6orf128 chromosome 6 open reading frame 128 884 CWF19L2 chromosome 20 open reading frame 70 FRKRA protein kinase, interferoninducible double strande 179 TGM7 transglutaminase 7 B3GALT7 UDPGal:betaGal beta 1,3galactosyltransferase polyp 74 FLJ10569 hypothetical protein FLJ10569 387 LOC197387 hypothetical LOC197387 600 C14orf68 chromosome 14 open reading frame 68 849 LOC400849 hypothetical LOC400849 799 FLJ32028 hypothetical protein FLJ32028 362 RBP7 retinol binding protein 7, cellular 201 FBI4 protein FLA domain family member 2 TEA domain family member 2	374907 55174 197387 283600 400849 201799 201799 116362 404201 8463 8463	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893 NP_689893 NP_443192 NP_995329 NP_003589 NP_003589	NM_205857 NM_003598 NM_003598 NM_024831	4D 4D	238323_at 238346_s_at
TEX13A TEX13A MGC50372 MGC20470 C6orf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7 B3GALT7 C14orf68 LOC197387 C14orf68 LOC400849 FLJ32028 RBP7 FB14 TEAD2 TEAD2	37490 55174 55174 19738 28360 400849 201799 11636 40420 8463	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893 NP_443192 NP_995329 NP_003589	NM_205857 NM_003598 NM_003598	4D	238323_at
TEX13A TEX13A MGC50372 MGC20470 C6orf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7 B3GALT7 C14orf68 LOC400849 FLJ32028 RBP7 FB14 TEAD2	37490 55174 19738 28360 40084 20179 11636 40420 8463	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893 NP_689893 NP_443192 NP_995329 NP_003589	NM_205857 NM_003598		220227 2 gt
TEX13A TEX13A MGC50372 MGC20470 C6orf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7 B3GALT7 C14orf68 LOC400849 FLJ32028 RBP7 FB14	37490 55174 55174 19738 283600 400844 201799 11636 40420	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893 NP_443192 NP_995329	NM_205857	1	1110111 2 2 4
TEX13A TEX13A MGC50372 MGC20470 C6orf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7 B3GALT7 C140rf68 LOC197387 C14orf68 LOC400849 FLJ32028 RBP7	37490 55174 55174 19738 28360 400849 201799	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893 NP_443192		4D	238081_at
TEX13A MGC50372 MGC20470 C6orf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7 B3GALT7 C14orf68 LOC400849 FLJ32028	37490 55174 19738 28360 400849 201799	NP_060612 XP_117030 NP_997000 XP_378988 NP_689893	NM 052960	4D	238066_at
TEX13A TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 3 C20orf70 PRKRA 9 TGM7 7 B3GALT7 7 B3GALT7 5 FLJ10569 7 LOC197387 0 C14orf68	37490° 55174 19738° 283600 400849	NP_060612 XP_117030 NP_997000 XP_378988	NM_152680	4D	238063_at
TEX13A TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 7 CTGM7 PRKRA 9 TGM7 B3GALT7 B3GALT7 C14orf68	37490° 55174 55174 19738° 283600	NP_060612 XP_117030 NP_997000	XM_378988	4D	237806_s_at
TEX13A TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 3 C20orf70 PRKRA 9 TGM7 7 B3GALT7 FLJ10569 FLJ10569	37490° 55174 19738°	NP_060612 XP_117030	NM_207117	4D	237765_at
3 IQCF2 TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 3 C20orf70 PRKRA 9 TGM7 7 B3GALT7	37490° 55174	NP_940942 NP_060612	XM_117030	4D	237730_at
TEX13A TEX13A MGC50372 MGC20470 Coorf128 CWF19L2 C20orf70 PRKRA TGM7 B3GALT7	37490	NP_940942	NM_018142	4D	237504_at
TEX13A TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 1 C20orf70 PRKRA 9 TGM7	37.00		NM_198540	40	20/000_41
TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2 3 C20orf70 PRKRA	116179	NP_443187	NM_052955	4D	237288_at
TEX13A TEX13A MGC50372 MGC20470 COorf128 CWF19L2 C20orf70	8575	NP_003681	NM_003690	4D	237107_at
TEX13A 3 MGC50372 0 MGC20470 8 C6orf128 4 CWF19L2	140683	NP_542141	NM_080574	4D	237099_at
TEX13A TEX13A MGC50372 MGC20470 C6orf128	143884	NP_689647	NM_152434	4D	237040_at
TEX13A 3 MGC50372 0 MGC20470	221468	NP_660359	NM_145316	4D	237016_at
TEX13A MGC50372	143630	NP_659490	NM_145053	4D	236965_at
TEX13A	253143	NP_775837	NM_173566	4D	236941_at
IQCF2	56157	NP_112564	NM_031274	4D	236848_s_at
TO COLD	389123	NP_982248	NM_203424	4D	236767_at
FLJ39485	285603	NP_787116	NM_175920	4D	236728_at
MDS009	56986	NP_064619	NM_020234	4D	236649_at
RPS27	6232	NP_001021	NM_001030	4D	236621_at
DOK6	220164	NP_689934	NM_152721	4D	236290_at
EIF3S9	8662	NP_003742	NM_003751	4D	236274_at
67 ZNF346 zinc finger protein 346	23567	NP_036411	NM_012279	4D	236267_at
MMRN2	79812	NP_079032	NM_024756	4D	236262_at
466 PHOSPHO1 phosphatase, orphan 1	162466	NP_848595	NM_178500	4D	236218_at
KIAA1853	84530	NP_919262	NM_194286	4D	236151_at
ABLIM2	84448	NP_115808	NM_032432	4D	236087_at
XAGE3	170626	NP_573440	NM_133179	4D	236040_at
392 C10orf78 chromosome 10 open reading frame 78	119392	NP_001002759	NM_001002759	4D	236027_at
tSNAREs					

insulin growth factorlike family member 1	IGFL1	374918	NP_940943	NM_198541	40	239430_at
HermanskyPudlak syndrome 1	HPS1	3257	NP_000186	11	4D	239382_at
hypothetical protein MGC11102	MGC11102	84285	NP_115701	NM_032325	4D	
hypothetical protein DKFZp547K1113	DKFZp547K111	374655	NP_940928	NM_198526	4D	239335_at
corin, serine protease	CORIN	10699	NP_006578	NM_006587	4D	239261_s_at
chromosome 6 open reading frame 93	C6orf93	84946	NP_116249	NM_032860	4D	239212_at
complement component (3b/4b) receptor 1, including	CRI	1378	NP_000564	NM_000573	4D	239205_s_at
ankyrin repeat domain 22	ANKRD22	118932	NP_653191	NM_144590	4D	239196_at
hypothetical protein MGC39372	MGC39372	221756	XP_376463	XM_376463	4D	239186_at
LOC440064	LOC440064	440064	XP_498525	XM_498525	4D	239081_at
hypothetical protein FLJ10534	FLJ10534	55720	NP_060598	NM_018128	4D	239042_at
aldolase A, fructosebisphosphate	ALDOA	226	NP_000025	NM_000034	4D	238996_x_at
densityregulated protein	DENR	8562	NP_003668	NM_003677	4D	238982_at
hypothetical protein FLJ38973	FLJ38973	205327	NP_710156	NM_153689	4D	238974_at
Kruppellike factor 12	KLF12	11278	NP_009180	NM_007249	4D	238940_at
TRAFinteracting protein with a forkheadassociated	TIFA	92610	NP_443096	NM_052864	40	238858_at
RIM binding protein 2	KIAA0318	23504	NP_056162	NM 015347	4D	238817 at
tyrosyltRNA synthetase	YARS	8565	NP_003671	NM_003680	4D	238760_at
Nacylphosphatidylethanolaminehydrolyzing phospholi	NAPEPLD	222236	NP_945341	NM_198990	4D	238722_x_at
hypothetical protein FLJ20582	FLJ20582	54989	NP_054825	NM_014106	4D	238687_x_at
WD repeat domain 36	WDR36	134430	NP_644810	NM_139281	4D	238677_at
chromosome 14 open reading frame 150	C14orf150	112840	NP_542397	NM_080666	4D	238650_x_at
hypothetical protein FLJ21657	FLJ21657	64417	NP_071928	NM_022483	4D	238635_at
hypothetical protein MGC2474	MGC2474	65988	NP_076420	NM_023931	4D	238606_at
Cblinteracting protein Sts1	STS1	84959	NP_116262	NM_032873	4D	238587_at
hypothetical protein MGC39389	FLJ32384	124790	NP_653209	NM_144608	4D	238547_at
synleurin	SLRN	353281	NP_852607	NM_181506	4D	238488_at
similar to PI3kinaserelated kinase SMG1 isoform I;	LOC440345	440345	XP_496125	XM_496125	4D	238449_at
citrate lyase beta like	CLYBL	171425	NP_996531	NM_206808	4D	238440_at
carbonic anhydrase VBlike	CA5BL	340591	XP_291346	XM_291346	4D	238435_at
hypothetical protein FLJ33069	FLJ33069	137835	NP_653250	NM_144649	4D	238429_at

zinc finger protein 214	ZNF214	1/61	NP_03/381	NW_013249	#5	24343/_5_41
zinc tinger protein 214	ZNF214	7/61	NP_03/381	NIM 013249	3 5	242450_at
hypothetical protein LOC339290	LOC339290	339290	XP_375527	XM_3/552/	£ £	243426_at
maba1	KIAA1324	57535	NP_065826	NM_020775	4D	243349_at
KIBRA protein	KIBRA	23286	NP_056053	NM_015238	4D	243290_at
FLJ35093 protein	FLJ35093	374986	NP_940951	NM_198549	4D	243042_at
ADPribosylation factorlike 8	ARL8	221079	NP_848930	NM_178815	4D	242727_at
hypothetical protein FLJ13305	FLJ13305	84140	XP_291019	XM_291019	4D	242584_at
TNF receptorassociated factor 4	TRAF4	9618	NP_004286	NM_004295	4D	242473_at
hypothetical protein LOC199899	LOC199899	199899	XP_378866	XM_378866	4D	242284_at
poly(A) polymerase beta (testis specific)	PAPOLB	56903	NP_064529	NM_020144	4D	242158_at
orthodenticle homolog 2 (Drosophila)	OTX2	5015	NP_068374	NM_021728	4D	242128_at
chromosome 6 open reading frame 152	C6orf152	167691	NP_859065	NM_181714	4D	242006_at
checkpoint suppressor 1	CHESI	1112	NP_005188	NM_005197	4D	241984_at
WD repeat domain 4	WDR4	10785	NP_387510	NM_033661	4D	241937_s_at
calmodulin binding transcription activator 1	CAMTAI	23261	NP_056030	NM_015215	4D	241882_at
sodium channel, nonvoltagegated 1, gamma	SCNNIG	6340	NP_001030	NM_001039	4D	241436_at
hypothetical protein FLJ34443	FLJ34443	285464	NP_787114	NM_175918	4D	241408_at
similar to hypothetical protein	LOC387816	387816	XP_370664	XM_370664	4D	241360_at
hypothetical LOC283034	LOC283034	283034	XP_210860	XM_210860	4D	241337_at
forkhead box N4	FOXN4	121643	NP_998761	NM_213596	4D	241009_at
keratin associated protein 193	KRTAP193	337970	NP_853640	NM_181609	4D	240967_at
fibroblast growth factor receptor 2 (bacteriaexpre	FGFR2	2263	NP_075262	NM_022973	4D	240913_at
similar to RIKEN cDNA 4933424N09 gene	MGC16943	112479	NP_542394	NM_080663	4D	240603_s_at
chromosome 20 open reading frame 71	C20orf71	128861	NP_848561	NM_178466	4D	240359_at
solute carrier family 15 (H+/peptide transporter).	SLC15A2	6565	NP_066568	NM_021082	4D	240159_at
hypothetical protein LOC201243	LOC201243	201243	NP_783861	NM_175734	4D	240082_s_at
lin7 homolog A (C. elegans)	LIN7A	8825	NP_004655	NM_004664	4D	240027_at
dachshund homolog 2 (Drosophila)	DACH2	117154	NP_444511	NM_053281	4D	239738_at
hypothetical protein MGC48972	MGC48972	123879	NP_775746	NM_173475	4D	239648_at
hypothetical protein MGC15912	MGC15912	84972	NP_116275	NM_032886	4D	239635_at
LOC440885	LOC440885	440885	XP_498906	XM_498906	4D	239624_at
hypothetical protein FLJ20481	FLJ20481	54947	NP_060309	NM_017839	4D	239598_s_at
hypothetical protein FLJ37659	FLJ37659	286499	NP_775969	NM_173698	4D	239481_at
zinc finger protein 284	ZNF284	342909	XP_496317	XM_496317	4D	239462_at

insuriogiucosamine surionydroiase (surramidase)	SUSII	0++0	TAT 000120	11111 000100		NY LOCATE IN
North Serie product	ecen.	6440	NOI NO DIN	NM 001000	45	35626 at
FI N29 gene product	FLN29	10906	NP 006691	NM 006700	4D	35254 at
reticulon 2	RTN2	6253	NP_005610	NM_005619	4D	34408_at
peroxisomal biogenesis factor 14	PEX14	5195	NP_004556	NM_004565	4D	33760_at
interferon stimulated gene 20kDa	ISG20	3669	NP_002192	NM_002201	4D	33304_at
myosin IC	MYOIC	4641	NP_203693	NM_033375	4D	32811_at
Nedd4 binding protein 1	N4BPI	9683	NP_694574	NM_153029	4D	32069_at
hypothetical protein BC002942	BC002942	91289	NP_149977	NM_033200	4D	31837_at
DEAD (AspGluAlaAsp) box polypeptide 49	DDX49	54555	NP_061943	NM_019070	4D	31807_at
bromodomain and WD repeat domain containing 3	BRWD3	254065	NP_694984	NM_153252	4D	244738_at
nuclear transcription factor Y, beta	NFYB	4801	NP_006157	NM_006166	4D	244704_at
hypothetical protein FLJ36674	FLJ36674	284040	NP_775893	NM_173622	4D	244698_at
cytochrome c, somatic	CYCS	54205	NP_061820	NM_018947	4D	244546_at
hypothetical LOC200624	LOC200624	200624	XP_117257	XM_117257	4D	244518_at
hypothetical protein MGC27466	MGC27466	127396	NP_689586	NM_152373	4D	244398_x_at
hypothetical protein FLJ31340	FLJ31340	222223	NP_689961	NM_152748	4D	244317_at
THAP domain containing 5	THAP5	168451	NP_872335	NM_182529	4D	244190_at
muskelin I, intracellular mediator containing kelc	MKLNI	4289	NP_037387	NM_013255	4D	244171_at
gammaaminobutyric acid (GABA) A receptor, alpha 1	GABRAI	2554	NP_000797	NM_000806	4D	244118_at
hypothetical protein FLJ35382	FLJ35382	163859	NP_689821	NM_152608	4D	244103_at
protein phosphatase 1K (PP2C domain containing)	PPMIK	152926	NP_689755	NM_152542	4D	244011_at
ATPbinding cassette, subfamily B (MDR/TAP), member	ABCB1	5243	NP_000918	NM_000927	4D	243951_at
Fraser syndrome 1	FRASI	80144	NP_079350	NM_025074	4D	243935_at
DKFZP434I116 protein	DKFZP4341116	25962	NP_056311	NM_015496	4D	243927_x_at
rab3 GTPaseactivating protein, noncatalytic subuni	RAB3GAP150	25782	NP_036546	NM_012414	4D	243851_at
collagen and calcium binding EGF domains 1	CCBEI	147372	NP_597716	NM_133459	4D	243805_at
similar to ZINC FINGER PROTEIN 257 (BONE MARROW ZI	MGC12518	90816	NP_291026	NM_033548	4D	243661_at
solute carrier family 1 (glutamate transporter), m	SLC1A7	6512	NP_006662	NM_006671	4D	243623_at
REV1like (yeast)	REVIL	51455	NP_057400	NM_016316	4D	243592_at
APG10 autophagy 10like (S. cerevisiae)	APG10L	83734	NP_113670	NM_031482	4D	243477_at
CasBrM (murine) ecotropic retroviral transforming	CBL	867	NP_005179	NM_005188	4D	243475_at

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signal transducer and activator of transcription 1	STAT1	6772		NP_009330	NM_007315	AFFX- HUMISGF3A/M 97934D
signal transducer and activator of transcription 1	STATI	6772		NP_009330	NM_007315	AFFX- HUMISGF3A/M 97934D
rabaptin, RAB GTPase binding effector protein 2	RABEP2	79874	NP_079092	NM_024816	4D	74694_s_at
amplified in breast cancer 1	ABC1	63897	NP_071353	NM_022070	4D	65493_at
apoptosisrelated protein PNAS1	FLJ39616	51275	NP_057618	NM_016534	4D	64432_at
SHQ1 homolog (S. cerevisiae)	SHQ1	55164	NP_060600	NM_018130	4D	63009_at
solute carrier family 24 (sodium/potassium/calcium	SLC24A3	57419	NP_065740	NM_020689	4D	57588_at
dishevelled, dsh homolog 2 (Drosophila)	DVL2	1856	NP_004413	NM_004422	4D	57532_at
hypothetical protein MGC13138	MGC13138	92595	NP_219363	NM_033410	4D	57516_at
KIAA1196 protein	KIAA1196	57473	NP_065764	NM_020713	4D	55872_at
chromosome 10 open reading frame 76	C10orf76	79591	NP_078817	NM_024541	4D	55662_at
HermanskyPudlak syndrome 4	HPS4	89781	NP_071364	NM_022081	4D	54037_at
RAN binding protein 10	RANBP10	57610	NP_065901	NM_020850	4D	53987_at
KIAA1698 protein	KIAA1698	80789	NP_085131	NM_030628	4D	53968_at
similar to RIKEN cDNA 3000004N20	LOC389677	389677	NP_976324	NM_203390	4D	51228_at
protein phosphatase methylesterase1	PMEI	51400	NP_057231	NM_016147	4D	49077_at
chromosome 5 open reading frame 4	C5orf4	10826	NP_115761	NM_032385	4D	48030_i_at
leucine zipper, putative tumor suppressor l	LZTS1	111178	NP_066300	NM_021020	4D	47550_at
abhydrolase domain containing 6	ABHD6	57406	NP_065727	NM_020676	4D	45288_at
gap junction protein, alpha 4, 37kDa (connexin 37)	GJA4	2701	NP_002051	NM_002060	4D	40687_at
DEAD (AspGluAlaAsp) box polypeptide 28	DDX28	55794	NP_060850	NM_018380	4D	40255_at
SH2B homolog	SH2B	25970	NP_056318	NM_015503	4D	40149_at
interleukin 1, beta	ILIB	3553	NP_000567	NM_000576	4D	39402_at
extra spindle poles like 1 (S. cerevisiae)	ESPL1	9700	NP_036423	NM_012291	4D	38158_at
parvin, beta	PARVB	29780	NP_001003828	NM_001003828	4D	37966_at
prolyl endopeptidase	PREP	5550	NP_002717	NM_002726	4D	37950_at
surfactant, pulmonaryassociated protein B	SFTPB	6439	NP_000533	NM_000542	4D	37004_at
vmaf musculoaponeurotic fibrosarcoma oncogene homo	MAFF	23764	NP_036455	NM_012323	4D	36711_at